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The present invention relates to new NOVX polypeptides. The invention is useful for treating or preventing a NOVX-associated disorder such as cardiomyopathy or atherosclerosis, where the disorder is related to cell signal processing and metabolic pathway modulation in a subject, signal processing and metabolic pathway modulation in a subject, or preferably human. The invention is also useful for treating metabolic disorders (e.g. obesity), diabetes, infectious disease, neurodegenerative disorders, haematopoietic disorders and various cancers. The molecules of the invention are also useful for treating or preventing cirrhosis. Cancertis, acne, hair growth, pigmentation disorders, endocrine disorders, respiratory disease, gastro-intestinal diseases, reproductive, health, neurological diseases, bone marrow transplantation, endocrine diseases, allergy and inflammation, nephrological disorders, urinary system disorders, neuropsychatric disorders and age-related disorders. The present nucleic acid sequence represents a PCR primer that was used in the methods of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gangolli EA, Patturajan M, Vernet CAM, Malyankar UM, Kekuda R;
Stone DJ, Anderson D, Shimkets RA, Burgess CE, Zerhusen BD, Liu X;
Spytek KA, Casman SJ, Boldog FL, Smithson G, Li L, Ji W;
neuropsychiatric disorder; EGF-related protein; SCUBE1; TEN-M4; adipocyte complement-related Clq tumour necrosis factor; out at first; beta adrenergic receptor kinase; Eph&G-Ghk-2; glucose transporter; type la membrane sushi-containing domain; butyrophilin; type la membrane-sushi domain containing; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated NOVX polypeptide useful for treating cardiomyopathy, atherosclerosis, metabolic disorders, diabetes, obesity, infectious disease, anorexia, neurodegenerative disorders, Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.0%; Score 13.8; DB 1; Length 18;
88.2%; Pred. No. 1.9e+02;
ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 3 A; 8 C; 5 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 198; 318pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      741
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-MAY-2001; 2001US-294075P.
24-JUL-2001; 2001US-307506P.
10-AUG-2001; 2001US-311590P.
10-AUG-2001; 2001US-311613P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                  2000US-257314P.
2001US-288153P.
                                                                                                                                                                                                                                                                                                                                                                                           2000US-265704P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001US-315617P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-SEP-2001; 2001US-322358P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   725 AGCAGGGGCCTGGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18 AGCATGGCGCCTGGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 88.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-590744/63.
                                                                                                                                                                                                                    WO200257453-A2
                                                                                                                                                                                                                                                                                                                            19-DEC-2001;
                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                            02-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-AUG-2001;
                                                                                                                                                                                                                                                                           25-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ27299
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ27299/
ID AAQ2
XX
AC AAQ2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a differential display comprising contacting cond, with a first primer that hybridises to an oligonucleic sequence coding for a leucine-rich motif, and with a second oligonucleotide primer to form a cDNA-hybrid molecule. The method comprises obtaining mRNA from at least 2 samples, synthesising CDNA from the RNA of each sample. Contacting the cDNA with a first primer that hybridises to an oligonucleotide primer to form cDNA-hybrid molecules, amplifying the oligonucleotide primer to form cDNA-hybrid molecules, amplifying the collgonucleotide primer to form cDNA-hybrid molecules, amplified products amplified products coding for at least one secreted or transmembrane protein. The method is useful for discovering novel secreted and/or transmembrane method is useful for discovering novel secreted and/or transmembrane collent are important for cell processes and play an important cole in determining its phenotype, and which act as mediators for the transfer of signals from external environment into the cell itself, thus modulating gene expression. Sequences ABX03792-ABX03869 represent DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cell signal processing, metabolic pathway modulation, metabolic disorder, obesity, diabetes; infectious disease, neurodegenerative disorder; acne; Alzheimer's disease; Parkinson's disease; immune disorder; cancer; haematcopoietic disorder; cirrhosis; pancreatitis; learning defect; memory defect; infertility; congenital heart defect; hair growth, pigmentation disorder; endocrine disorder; respiratory disease; health; gastro-intestinal disease; reproductive; neurological disease; health; bone marrow transplantation; endocrine disease; allergy; inflammation; nephrological disorder; urinary system disorder; age-related disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Differential display method for identifying secreted or transmembrane protein, comprises contacting a DNA with a first primer that hybridizes to a sequence coding for a leucine-rich motif and with a second
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; NOVX; NOVX-associated disorder; cardiomyopathy; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
1.0%; Score 13.8; DB 1; Length 18;
Best Local Similarity 88.2%; Pred. No. 1.96+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoding secreted protein signal peptide sequences.
                                                                                                                                                                                                                                                                     (UYRA-) UNIV RAMOT APPLIED RES & IND DEV LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 1 A; 8 C; 5 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Fig 2; 37pp; English.
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                                                                                                                                                           23-JAN-2002; 2002WO-IL00071.
                                                                                                                                                                                                                 23-JAN-2001; 2001US-263158P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        566 CACTGCTCCAGCAGGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CACTGCTCCTGCTGGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-NOV-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligonucleotide primer
                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-599769/64.
                                                                                                                                                                                                                                                                                                                                                                                                            P-PSDB; ABG98328
                                                   WO200259259-A2
                                                                                                                                                                                                                                                                                                                            Wreschner DH;
Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABS52111;
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Gaps

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AAQ47696 standard; DNA; 19
                                                                        Buckler AJ, Chang DD,
                                                                                                                                                                                           Best Local Similarity
                                                                                WPI; 1992-284656/34.
             pSPL1 primer DHA14.
                                                           28-JAN-1991;
                                      WO9213071-A1
                                                    27-JAN-1992;
   25-MAR-2003
02-FEB-1993
                                                                                                                                                                                                                                             25-MAR-2003
04-FEB-1994
                                             06-AUG-1992.
                               Synthetic.
                                                                                                                                                                                                                                      AAQ47696;
                                                                                                                                                                                                             11
                                                                                                                                                                                       Query Match
                                                                                                                                                                                               Matches
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RNaseH; RNA cleavage; DNA cleavage; hybridisation; protein kinase C gene; gene expression modulation; ras; raf; therapy; AIDS; atherosclerosis; infection; cell growth; ss.
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                                                                                                                                                                                                                                                                                                                              Cyclophilin C-associated membrane proteins and DNA - used for screening for immunomodulatory agents and for diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                        Primers CAMP-c #2 and CAMP-c #4 were used to screen a plasmid cDNA library derived from AC 6 cells for CDNA clones encoding CAMP-c. These primers amplify a fragment of approx. 250 bp. (Updated on 25-MAR-2003 to correct PN field.)
Cyclophilin associated membrane protein; CAMP-c; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.0%; Score 13.8; DB 1; Length 19;
88.2%; Pred. No. 2e+02;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Target sequence for 2'-substituted oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 3 A; 6 C; 6 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monia
                                                                                                                                                                                                                  (STRD ) UNIV LELAND STANFORD JUNIOR.
                                                                                                                                                                                                                                                                                                                                                                                                         Example; Page 64; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Martin P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ٣
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97WO-US02043.
                                                                                                                                              93WO-US01123
                                                                                                                                                                                   92US-0832862
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  235 CAGGCATCTGCATCTGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX36444 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cook PD,
                                                                                                                                                                                                                                                                                             WPI; 1993-272887/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM I (NOVS ) NOVARTIS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1997-424968/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
                                                                                                                                              08-FEB-1993;
                                                                                                                                                                               07-FEB-1992;
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                                                                     WO9316183-A1
                                                                                                                                                                                                                                                        Friedman JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-JUL-1999
                                                                                                          19-AUG-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Altmann K,
                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX36444;
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                                                                                                                                                                                                                                                                                                                                                                    therapy
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AAX36444/C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fragments of a mouse cosmid clone, Malphacd#9, known to contain
exon sequences of the Na, R-ATFase alphal subunit gene (Tam, S.-
Y., et al., Mol. Cell. Biol. 10:6619-6623 (1990) were used to
demonstrate that when a fragment contg. an entire exon with
flanking intron sequences (in the proper orientation) is inserted
into an intron of an in vivo splicing plasmid, the exon is retained
in the mature poly A+ cytoplasmic RNA. In vivo splicing plasmid
DSPLI was used. The insertion site is within an intron from the
HIV-1 tat gene whose flanking exons and splice sites were substituted
for the second intron of the rabbit beta-q-lobin gene.
Oligonucleotide pairs for PCR are represented in AAQ27300-01).
The antisense primers DHAB14 and SA2 (AAQ27299 and AAQ27301) were used
as primer in the first strand cDNA synthesis reactions.
SDI and SA1 (AAQ27302-03) are internal to the initial RNA/PCR prod.
and were used for reamplification of RNA/PCR prods.
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                                                                                                  Cosmid; MalphaG#9; exon; Na, K-ATPase; in vivo splicing splasmid; HIV-1 tat; splice site; beta-globin; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolating coding sequence from mammalian genomic DNA - by providing fragmented DNA, inserting DNA into intron of in vivo splicing plasmid, introducing obtd. in vivo splicing plasmid construct into host cell, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence of primer CAMP-c #2 for cyclophilin associated membrane protein (CAMP-c) cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 13.8; DB 1;
Pred. No. 2e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                Sharp PA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 3 A; 5 C; 8 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                         (MASI ) MASSACHUSETTS INST TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                Housman DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 24; Page 33; 38pp; English.
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88.2%;
                                                                                                                                                                                                                                                                  92WO-US00692
                                                                                                                                                                                                                                                                                                       91US-0646664
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           (updated)
(first entry)
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15; Conservative
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Gaps

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2'-0-CH2-CH2-O-CH3 and 2'-deoxy sugar moieties, useful for therapy

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The invention relates to oligonucleotides (A), which specifically chiefless to RNA or Diagonucleotides (A), which specifically hybridises to RNA or DNA, comprises a linear sequence of nucleotide units linked by phosphodiseter or phosphorothicate linkages, comprising a first subsequence having 2'-0-H2-CH2-O-CH3 organ moieties and a second cultural accordance having 2'-deoxy sugar moieties. (A), which has RNASSH cativity for leaving a complementary strand, can be used to modulate the expression of ras, raf and protein kinase C genes, useful in the therapy of aberrant cell growth in humans, animals or plants. (A) can also be used diagnostically, particularly when labelled, to detect overexpression of mRNA or expression of abnormal RNA, including imaging of tissue sections, and as a research reagent. (A) has increased binding affinity for complementary strands (attributable to the 2'-O-CH2-CH2-O-CH3 sugar links), and increased resistance to nuclease (from the modified links and the collections).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; adenomatous Polyposis coli; APC; diagnosis; prognosis; neoplastic tissue; tumour tissue; tumour repressor; mutation; sporadic colorectal cancer; detection; PCR primer; SRP19; SSCP; single stranded conformation polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hedge PJ;
Thliveris A;
                                                                         This sequence represents a target sequence used to test the oligonucleotides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                          Score 13.8; DB 1; Length 19;
Pred. No. 2e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Groden J,
Nakamura Y,
                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 13 A; 4 C; 2 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                               the 2'-0-CH2-CH2-0-CH3 sugar moiety).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Carlson M,
Markham A,
                                          Example 2; Page 21; 86pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Primer for SSCP analysis of SRP19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JAPANESE FOUND CANCER RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   1141 GCCTTTTTTTTTTT 1157
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                                                                                                                                                                                                                                                                                                                                                                                            1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19 GCGTTTTTTTTTTTG 3
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91GB-0000974.
91GB-0000975.
91US-0741940.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (updated)
(first entry)
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UNIV UTAH.
                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Albertsen H, Anand R
Joslyn G, Kinzler K,
                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
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               or diagnosis
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                                                                                                                                                                                                                                                                                                                                                                                                                       15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-JAN-1991;
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11-MAR-1998
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(UTAH)
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Cancer diagnosis - by detecting mutation(s) in adenomatous polyposis
                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                        The present sequence is a primer for the SSCP analysis of SRP19, which was used in the development of a novel method of diagnosing or prognosing a human adenomatous Polyposis coli (APC) gene associated neoplastic tissue. The method comprises comparing APC gene coding sequences or mRNA in a tumour tissue, to APC gene coding sequences or mRNA in a non-neoplastic tissue, where a difference indicates an APC gene associated neoplasia of the tumour tissue. APC is a tumour repressor expressed in most normal tissues. APC mutations are found in familial adenomatous polyposis and sporadic colorectal cancer patients. The method enables mutations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New Alzheimer's disease related gene, AD4 - used to develop prods.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schellenberg GD;
                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Autosomal dominant early-onset Alzheimer's Disease; AD4; STM2; neurodegeneration; senile dementia; human chromosome 1; Volga German kindred; VG; yeast artificial chromosome library; expressed sequence tag database; polymerase chain reaction; PCR primer; Homo sanolens: sa.
                                                                                                                                                                                                                                                                     to be detected to provide an indication of predisposition to
                                                                                                                                                                                                                                                                                                                                                             1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2e+02; Ive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                    (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 0 A; 4 C; 8 G; 7 T; 0 other;
                                                                                                Example 8; Columns 31-32; 140pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human AD4 gene PCR primer INTIR.
                                                                                                                                                                                                                                                                                                                                                                                                                    100 ACAACCCCGGAGGCGCA 116
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                                                                                                                                                                                                                                                                                                                                                                            88.28;
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95US-0001675.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                17 ACAACCCAGGAGCCGCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 88.2
Matches 15; Conservative
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VA MEDICAL CENT.
 RL;
 White
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Wasco W;
                            WPI; 1997-372053/34.
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Vogelstein B,
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28-JUL-1995;
11-AUG-1995;
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                                                                        coli gene
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Tanzi RE,
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DP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphic marker; allele-specific; probe; amplification; PCR primer; hybridisation; plant; hybrid certification; genetic contribution; progeny; back-cross; hybrid; ancestry; corn; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                           A genetically isolated group of families with autosomal dominant early-onset Alzheimer's Disease (AD) has been studied and initial mapping analyses have predicted the AD4 locus (also known as STWA) resides on chromosome 1. The present sequence corresponds to a PCR primer which was used during the cloning procedure to isolate and sequence the AD4 gene. The group of families has been designated the Volga German (VG) kindreds. The entire gene has been amplified from VG individuals and unaffected individuals (from VG and unrelated lineages). Sequence analysis has shown that affected individuals have a nucleotide change at codon 141 resulting in an especially one in which Asn at position 141 has been replaced by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Brassica species allele-specific oligonucleotide probes and primers - useful for plant breeding
                                                                                                                                                                                                                                                                              Ile, can be used in a peptide vaccine. Detection of mutant AD4, for example using antibodies specific for the protein or using nucleic acid probes specific for the mutant gene, provides a means of
for detecting pre-disposition to or for diagnosis, prevention or treatment of Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                   Score 13.8; DB 1; Length 19;
Pred. No. 2e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Indels
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                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 6 A; 2 C; 10 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zea mays genome reverse PCR primer #274.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Murigneux A,
                                               Disclosure, Fig 11; 83pp; English.
                                                                                                                                                                                                                                                                                                                                   diagnosing Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    581 CCCTCCGTCTGCCCCC 597
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88.2%;
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96US-0032069
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lemieux B,
                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, to monitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a back-crossed plant with an ancestral plant.
                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphic marker, allele-specific, probe; amplification; PCR primer, hybridisation; plant; hybrid certification; genetic contribution; progeny; back-cross; hybrid; ancestry; corn; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV51705-V52008 are reverse PCR primers used to amplify fragments of th Zea mays genome in order to detect polymorphic markers. Such markers cabe used in the construction of allele-specific primers and probes for amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, to monitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a back-crossed plant with an ancestral plant.
                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Brassica species allele-specific oligonucleotide probes and primers - useful for plant breeding
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                                                                                                                         Score 13.8; DB 1; Length 19; Pred. No. 2e+02;
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                                                                                                                                                        Indels
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18.2%; Pred. No. 2e+02;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sapolsky RJ;
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                                                                                           Sequence 19 BP; 5 A; 6 C; 7 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 5 A; 6 C; 7 G; 1 T; 0 other;
                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                               Zea mays genome reverse PCR primer #275.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murigneux A,
                                                                                                                                                                                     568
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                                                                                                                                                                                                                                                                                          BP.
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                                                                                                                         1.0%;
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96US-0032069
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                                                                                                                                                                                   552 GGCAGGCATGCACACAC
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Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zea mays.
                                                                                                                                                                                                                                                                                                                      AAV51979;
                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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Matches
                                                                                                                                                                                                                                                          RESULT 283
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    888888
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Human; adenomatous Polyposis coli; APC; diagnosis; prognosis; neoplastic tissue; tumour repsressor; mutation; sporadic colorectal cancer; detection; PCR primer; SRP19; SSCP; single stranded conformation polymorphism; ss.

Homo sapiens

Synthetic.

US5691454-A.

Primer for SSCP analysis of SRP19.

(updated)
(first entry)

25-MAR-2003 08-APR-1998

AAT96201;

AAT96201 standard; DNA; 19 BP.

us09904568-3.rng

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AAT96201/c
         AAV56477-V56581 are primers used in the isolation of a human familial adenomatous polyposis coli (APC) protein from clone DP2.5. The gene for the protein is present on human chromsome $241 and is also referred to as adenomatous polyposis coli gene. It is a tumour suppressor gene, and mutations in this gene have been associated with tumourigenesis in retinoblastoma and colorectal tumours, and especially familial adenomatous polyposis (RAP) and Gardner's Syndrome (GS). The protein can be used in therapy to replace lack of native functional protein and the nucleic acids can be used for gene therapy. The nucleic acids that encode them can also be used as probes and primers in detection of the cancers and predisposition to it.
(Updated on 25-MAR-2003 to correct PR field.)
                                                                                                          Familial adenomatous polyposis coli, APC, tumour suppressor; therapy, chromosome 5q21; tumourigenesis; retinoblastoma; colorectal tumour; FAP; Gardner's Syndrome; GS; predisposition; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                        Hedge PJ;
, Thliveris A;
                                                                                                                                                                                                                                                                                                                                                                                                              Adenomatous polyposis coli protein - useful in the treatment cancers associated with mutation(s) on human chromosome 5g21
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Nakamura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 0 A; 4 C; 8 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 8; Column 31-32; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                      Anand R, Carlson M,
nzler K, Markham AF,
                         AAV56488 standard; DNA; 19 BP.
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91GB-0000963.
91GB-0000974.
91GB-0000975.
                                                                                                                                                                                                                   95US-0452655
                                                                                                                                                                                                                                                                                    94US-0289548
                                                                                                                                                                                                                                                                                                                UNIV JOHNS HOPKINS. UNIV UTAH.
                                                                                          Human DP2.5 APC primer #11
                                                              (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 88.2.
Best Local Si Conservative
                                                                                                                                                                                                                                                                                                                                                                          White RL;
                                                                                                                                                                                                                                                                                                                                                               Joslyn G, Kinzler K,
Vogelstein B, White
                                                                                                                                                                                                                                                                                                                                   ZENECA PHARM.
                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-427100/36.
                                                                                                                                                                                                                                                                                                       CANCER INST
                                                                                                                                                                                                                                                                                                                                                      Albertsen H,
                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                          08-AUG-1991;
                                                                                                                                                                                                                   25-MAY-1995;
                                                                                                                                                                                                                                                         16-JAN-1991;
                                                                                                                                                                                                                                                                16-JAN-1991;
                                                                                                                                                                                                                                      16-JAN-1991;
16-JAN-1991;
                                                                                                                                                                              US5783666-A.
                                                              25-MAR-2003
23-NOV-1998
                                                                                                                                                                                                21-JUL-1998
                                                                                                                                                 Synthetic
                                           AAV56488;
                                                                                                                                                                                                                                                                                                               (UYJO )
(UTAH )
(ZENE )
                                                                                                                                                                                                                                                                                                       CANC-)
       RESULT 284
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                                                         The present sequence is a primer for the SSCP analysis of SRP19, which was used in the development of a novel method of diagnosing or prognosing a human adenomatous Polyposis coli (APC) gene associated neoplastic fissue. The method comprises comparing APC gene coding sequences or mRNA in a tumour tissue, to APC gene coding sequences or mRNA in a non-neoplastic tissue, where a difference indicates an APC gene associated neoplasis of the tumour tissue. APC is a tumour repressor expressed in most normal tissues APC mutations are found in familial adenomatous polyposis and sporadic colorectal cancer patients. The method enables mutations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
.
                                                                                                                                                                                                                                                                                                                                                                                                                   to be detected to provide an indication of predisposition to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.0%; Score 13.8; DB 1; Length 19; 38.2%; Pred. No. 2e+02; lve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 0 A; 4 C; 8 G; 7 T; 0 other;
Example 8; Columns 25-26; 107pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100 ACAACCCCGGAGGCGCA 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Acadeccadadeccada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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Antibodies to normal and mutant adenomatous polyposis coli proteins - useful for detecting genetic predisposition to cancer

Groden J, Hedge PJ; Nakamura Y, Thliveris A;

Carlson M, Markham AF,

Albertsen H, Anand R, (Joslyn G, Kinzler K, M Vogelstein B, White RL;

WPI; 1998-017712/02.

IMPERIAL CHEM IND PLC. UNIV JOHNS HOPKINS. UNIV UTAH.

(ICIL ) (UYJO ) I (UTAH ) (CANC-)

CANCER INST

91GB-0000963. 91GB-0000974. 91GB-0000975. 91US-0741940. 94US-0289548.

16-JAN-1991; 16-JAN-1991; 16-JAN-1991; 08-AUG-1991; 12-AUG-1994;

95US-0452654 91GB-0000962

25-MAY-1995;

.6-JAN-1991

25-NOV-1997

g

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ö

Gaps

0

1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2e+02; Live 0; Mismatches 2; Indels

100 ACAACCCCGGAGGCGCA 116

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ACAACCCAGGAGCCGCA 1

17

RESULT 285

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The present sequence represents a primer for the wheat viviparous 1 (VPI) gene, which keeps mature seeds dormant and inhibits germination. The present invention describes genes which are homologues of the maize Viviparous 1 gene, obtained from oat Avena fatua and wheat which encode bolypeptides designated afVPI and taVPI respectively. The VPI activity keeps mature seeds dormant and inhibits germination and can be used to maintain or impose sufficient intensity and diration of dormancy to avoid pre-harvest sprouting (PHS) before harvest. The products can be used in the production of transformed crop plants having desirable primary or secondary dormancy. or after-ripening properties, and in particular may be resistant to PHS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified; deoxyfuranosyl moiety; therapy; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 19;
                                                                                                                                                                                                                                                                                  New isolated oat and wheat VPl genes, used, e.g. to impose sufficient dormancy to avoid pre-harvest sprouting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2' modified oligo used in the course of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 13.8; DB 1;
Pred. No. 2e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 3 A; 4 C; 11 G; 1 T; 0 other;
                                                                                                                                                                                                                   Holdsworth MJ;
                                                                                                                                                                                                                                                                                                                                    Claim 56; Page 89; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AIDS; atherosclerosis; tumour; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  800 CTCGCTCCCTGCAGCCG 816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.0%;
                                                                                                                                                                               (PLAN-) PLANT BIOSCIENCE LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90US-0463358
90US-0566977.
91WO-US05720.
92US-0835932.
92US-0854634.
                                                                                                               98WO-GB02835
                                                                                                                                               97GB-0020060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95US-0468037.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95US-0468037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18 CTCGCACCCTGCTGCCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX05488 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity 88.2
                                                                                                                                                                                                                Gale MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                  WPI; 1999-244424/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nuclease resistant;
          Triticum aestivum.
                                           WO9915667-A1
                                                                                                                                                                                                                Flintham JE,
                                                                                                             18-SEP-1998;
                                                                                                                                             19-SEP-1997;
                                                                             01-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US5859221-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-AUG-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX05488;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX05488/c
          à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 qq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a phosphorothioate antisense oligonucleotide which a gapped 2' modified oligonucleotide, whereby one part has at least two consecutive 2-F (2-H) nucleotides and the second part has at least five consecutive nucleotides with 2'-H sugar moieties. The modified oligonucleotide has increased nuclease resistance, and increased binding affinity for substrates. The oligonucleotide elicits RNase H strand cleavage of specific RNas oligonucleotide elicits RNase H strand cleavage of specific RNas oligonucleotide conditions as succeptible to oligonucleotide therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                   Antisense oligonucleotide, nuclease resistance, RNAse H strand cleavage;
phosphorothicate; oligonucleotide therapeutic; AIDS; atherosclerosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wheat; oat; viviparous 1; VPI; afVPI; taVPI; maize; detection; PHS; pre-harvest sprouting; dormant; germination; crop plant; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonuclectides eliciting RNAase H activity useful for diagnosis and treatment of diseases e.g AIDS or atherosclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ·,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13.8; DB 1; Length 19;
Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                   Antisense oligonucleotide which is gapped 2' modified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (Updated on 20-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 13 A; 4 C; 2 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wheat viviparous 1 (taVP1) primer #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Column 13; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (e.g. AIDS and atherosclerosis)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
AAZ10311/c
ID AAZ10311 standard; DNA; 19 BP.
XX
                                                                                                                                                                                                                                                                                                                                                          91US-0814961.
92WO-US11339.
94US-0244993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ന
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%;
                                                                                                                                                                                                                                                                                                                             95US-0465880.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCGTTTTTTTTTTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX33236 standard; DNA; 19
                                                                                    (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1141 GCCTTTTTTTCTTT
                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-539598/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Monia BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                          24-DEC-1991;
                                                                                                                                                                                                                                                                                                                           06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                              23-DEC-1992;
21-JUN-1994;
                                                                                  20-MAR-2003
08-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JUN-1999
                                                                                                                                                                                                                                                     USS955589-A.
                                                                                                                                                                                                                                                                                         21-SEP-1999
                                                                                                                                                                                                                       Synthetic.
                                                  AAZ10311;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cook PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13
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AAX33236/

ð d 0

Gaps .. 0 91GB-0000975. 91US-0741940. 94US-0289548.

16-JAN-1991; 08-AUG-1991; 12-AUG-1994;

Kawasaki AM;

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Cook PD,
                       Matches
                           RESULT 289
                            AAA93490/c
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The invention relates to a nuclease resistant compound that hybridises with RNA or DNA. The compound comprises covalently-bound nucleosides that individually include a ribose or deoxyribose sugar portion and a base portion, where the nucleosides are joined together by nucleosides internucleoside linkages such that is complementary to a RNA base sequence form a mixed base sequence; and where at least 1 of the nucleosides or to a DNA base sequence; and where at least 1 of the nucleosides includes a modified deoxyfuranosyl moiety bearing a 2'-substituent selected from cyano, fluoromethyl, thioalkoxyl, alkylsulphinyl, allyloxy and alkeneoxy groups. The nuclease resistant class alkylsulphonyl, allyloxy and alkeneoxy groups. The nuclease resistant close used for treating organisms having a disease characterised by the undesired production of a protein. They can be used in therapeutic cor prophylactic treatment in organisms such as bacteria, yeast, protozoa algae, plant and higher animal forms inflections and diagnosis. The ONS provide enhanced binding to terresting inflections and diagnosis. The ONS provide sequence-specific ONS provides superior potency and specificity compared to phosphorus-modified ons the present sequence represents a compared to phosphorus-modified ons the present sequence represents a construction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                               Nuclease resistant oligonucleotide analogues - having nucleosides including modified deoxyfuranosyl moiety bearing 2'-substituent to increase binding affinity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 13 A; 4 C; 2 G; 0 U; 0 other;
                                                                                                                                                                                                                                                           Example 19; Column 40; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
WPI; 1999-120005/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15;
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AAA84760;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 290
AAA84760
                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             셤
                                                                                                                                                                                                                                                                                                                                                                                    SRP19 gene; signal recognition particle protein; human; chromosome 5q21; familial adenomatous polyposis; FAP locus; Gardner's syndrome; GS; goradic tumour; adenoma; carcinoma; cancer; lung; breat; colon; rectum; bladder; liver; sarcoma; stomach; prostate; leukaemia; lymphoma; tumour suppressor; anti-APC antibody; detection; diagnosis; prognosis; genetic predisposition; drug screening; Adenomatous Polyposis Coli;
                                                                                                                                                                                                                                                                                                                                           Human SRP19 gene exon 1 PCR primer 1.
1141 GCCTTTTTTTTTTT 1157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APC gene; exon; PCR primer; ss.
                                                                                                                                                                                  AAA93490 standard; DNA; 19 BP.
                                               scerririririris 3
                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                      16-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US6114124-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-SEP-2000
                                                                                                                                                                                                                                    AAA93490;
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91GB-0000962. 91GB-0000963. 91GB-0000974.

16-JAN-1991; 16-JAN-1991; 16-JAN-1991;

95US-0450582

25-MAY-1995;

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The invention relates to a novel method for detecting Adenomatous Polyposis Coli (APC) protein in a sample. The method involves contexting the sample with antibodies which specifically binds to the contexting the sample with hutilibodies which specifically binds to the protein, and detecting an APC-antibody complex. Mutations in the APC gene play a role in tumorigenesis, indicating that it is a tumour contemplate of the APC gene play a role in tumorigenesis, indicating that it is a tumour the FAP (familial adenomatous polyposis) locus. FAP is an autosomal dominant inherited disease in which affected individuals developm to the FAP (familial adenomatous polyposis) locus. FAP is an autosomal dominant inherited disease in which affected individuals developm to some of which progress to malignancy. The FAP locus is often found to some of which progress to malignancy. The FAP locus is often found to be deleted in sporadic (i.e., non-familial) adenomas and carcinomas, and chromosome 5g deletions have also been observed in tumours of the lung, breast, colon, rectum, bladder, liver, sarcomas, stomach, and prostate, and in leukaemias and lymphomas. Although the FAP locus contains of several other genes such as FBK, TBJ, TBZ, and MCC, it is thought that mutations in the APC gene play a key role in the development of FAP and sporadic tumours. The method is useful for detecting APC protein and its mutations in the APC gene play a placental tissue, aministic fullid, blood, cancer, for detecting germiline and somatic alteration of will drype APC genes, and for testing therapeutic agents for the ability to suppress tumours. Sequences AAA93490-A93499 represent PCR primers used in an example in the APC and primer way in a normal processing the averagion of the human contains approached by a sequence of the human contains of the human contains of the human contains apple appressing to a normal processing the averagion of the human contains apple appressing to a normal processing to a normal processing to a normal processing to a
                                                                                                                                                                                                                                                                                                                                                               Detecting Adenomatous Polyposis Coli (APC) protein in a sample for diagnosing cancers, involves contacting the sample with antibodies that specifically bind to APC protein and detecting the complex formed -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                          Anand R;
                                                                                                                                                                                                                       Joslyn G, Kinzler K, Markham AF, Anand F, Thliveris A, Nakamura Y, Vogelstein B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2e+02; 2; Indels tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ribozyme; hairpin; hammerhead; gene therapy; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SRP19 (signal recognition particle protein) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 0 A; 4 C; 8 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 8; Column 31; 125pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cyclin F ribozyme binding site #28.
                                                                                            (ICIL ) IMPERIAL CHEM IND PLC. (UVJO ) UNIV JOHNS HOPKINS. (UTAH ) UNIV UTAH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100 ACAACCCCGGAGGCGCA 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17 ACAACCCAGGAGCCGCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA84760 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-DEC-2000 (first entry)
                                                                                                                                                                                                                  Groden J, J,
White RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15; Conservative
                                                                                                                                                                    (CANC-) CANCER INST.
                                                                                                                                                                                                                                                                                                                   WPI; 2000-565003/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        restenosis; ss.
                                                                                                                                                                                                                                            Albertsen H,
                                                                                                                                                                                                                    Carlson M,
                                                                                                                                                                                                                                                                     Hedge PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mammalia
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17-FEB-2000,

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The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinases other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1.
Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in restenosis treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag= c
/note= "Contains a 2'-0-substituent selected from
2'-0-aminooxyethyl, 2'-0-ethylaminooxyethyl and
2'-0-dimethylaminooxyethyl"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Contains a 2'-0-substituent selected from 2'-0-aminooxyethyl, 2'-0-ethylaminooxyethyl and 2'-0-dimethylaminooxyethyl"
                                                                                                                                                                                           New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDKI, PCNA and Cyclin B1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Phosphorothioate; modified nucleoside; 2'-substituted; 2'-deoxy-erythro-pentofuranosyl sugar moiety; nuclease resistant; hybridisation; binding affinity; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                    1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              a
"Phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                             2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Phosphorothioate oligonucleotide, SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 3 A; 6 C; 6 G; 4 T; 0 other;
                                                                                                                                             Robbins JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                               Disclosure; Page 82; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   location/Qualifiers
                                                                                                                                             Barber JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA06830 standard; DNA; 19 BP
                                                                99WO-US28772.
                                                                                         98US-0110954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        877 GCCAAGTTCCAGGAGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 geckgerreckggkger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                              15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1..3
/*tag=
/note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18..19
                                                                                                                                         Welch PJ,
                                                                                                                                                                    WPI; 2000-412314/35.
                                                                                                                (IMMI-) IMMISOL INC
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sest Local Similarity
          WO200032765-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200008042-A1
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
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                                                                                         04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-JUN-2000
                                                             06-DEC-1999;
                                   08-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA06830;
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                           Tritz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 291
                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA06830,
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oligonuclectides and to oligonuclectides that elicit RNase H for cleavage in a complementary nucleic acid strand. It also relates to cleavage in a complementary nucleic acid strand. It also relates to cleavage in a complementary nucleic acid strand. It also relates to cligonuclectides wherein at least some of the nucleotides include a substituent that potentiates hybridisation of the nucleotides include a 2'-decxy-exythro-pentofurancyl sugar moiety. The nucleotides include a 2'-decxy-exythro-pentofurancyl sugar moiety. The inclusion of one or more aminoxy moieties in such oligonucleotides to a complementary strand. The oligonucleotides to a complementary strand. The oligonucleotides of the invention are used as diagnostic, therapeutic or research reagents, and can be used to modulate modified nucleosides have increased nuclease resistance and increased
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                represents a phosphorothioate oligonucleotide used in an exemplification of the present invention which has 2'-substituted regions flanking a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             blood disorder, hasmophilia, autoimmune disease, diabetes, inflammation, infection, fungal, bacterial, viral, HIV, allergy, arthritis, neurodegenerative disease, asthma; contraceptive; hybridisation probe,
                                                                                                                                                                                                                                   Aminooxy-modified nucleosides and oligonucleotides useful in diagnostic, therapeutic and research reagents and for modulating the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; secreted protein; cancer; tumour; cardiovascular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              binding affinity to a complementary strand. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    central region. This sequence is the complement of AAA06829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                 The invention relates to aminooxy-modified nucleosides and
                                                                                                                                                                  Prakash TP, Kawasaki AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Clone vc46_1 hybridisation probe, SEQ ID NO:96.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 13 A; 4 C; 2 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                               Example 40; Page 75; 195pp; English.
                                                                                                                                                                                                                                                                          expression of protein in organisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1141 GCCTTTTTTTTTTT 1157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA23478 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19 GCGriririririris 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99WO-US19351.
                                                      99WO-US17988
                                                                                          98US-0130973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                Cook PD,
                                                                                                                             PHARM INC
                                                                                                                                                                                                    WPI; 2000-224020/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200011015-A1
                                                                                                                           SISI (-SISI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                    09-AUG-1999;
                                                                                        07-AUG-1998;
                                                                                                                                                                Manoharan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.9-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA23478;
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Gaps

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WO200014198-A2 Homo sapiens,

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The invention relates to 40 human secreted proteins (AAV94981-V95020),
and cDNA sequences encoding them (AAA23423-A23462). The secreted
control of the invention include those that are thought to be only
proteins of the invention include those that are thought to be only
partially secreted, i.e., transmembrane proteins. The proteins of the
corporation may exhibit one or more activities selected from the following:
corporation activity, cell proliferation; differentiation; immune
modulation; haematopoiesis regulation; tissue growth activity;
condition activity; anti-inflammatory activity; and tumour
activin/inhibin activity; anti-inflammatory activity; and tumour
condition activity. The proteins may be administered to patients as
condition activity anti-inflammatory activity; and tumour
condition activity. The proteins may be used as part of a gene therapy
concludes of the invention include autoimmune diseases; genetic
disorders, haemophila; cardiovascular diseases; cancer; bacterial,
concluders, haemophila; cardiovascular diseases; denetic
disorders, haemophila; cardiovascular diseases; bacterial,
concluders, cardiovascular diseases; denetic
disorders, haemophila; cardiovascular diseases, denetic
constituin dependent diabetes mellitus; and allergic reactions such as
concluders, osteoporosis, osteoarthritis, periodontal diseases, Alzheimer's
concluders, coreoporosis, osteoarthritis, periodontal diseases, Alzheimer's
concluders, pervention may also be used for treating wounds, burns,
disease, parkinson's disease, Huntington's disease and amyoricophic
conditionally be useful as contraceptives. Nucleic acid sequences of the
converse and northers and nuches Reminence and sequences of the
converse and nuches and nuches Reminence and represent
                                                                                                                                                                                                                                                                                                                                    secreted or transmembrane proteins and polynucleotides encoding n, useful for treating neurodegenerative disorders, autoimmune
                                                                                                                                                                                                                                                      Hall J, Rapiejko P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 5 A; 4 C; 7 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 340; 357pp; English.
                                                                                                                                                                                                                                                    Hoffman H,
  98US-0097638.
98US-0097659.
98US-0099618.
                                                               98US-0102092.
98US-0109978.
98US-0113645.
                                                                                                                                   98US-0113646
                                                                                                                                                                                                                                             Valenzuela D, Yuan O,
                                                                                                                                                                                                     (ALPH-) ALPHAGENE INC.
                                                                                                                                                                                                                                                                                                                                                                                   diseases and cancer -
                                                                                                                                                                                                                                                                                          WPI; 2000-224657/19.
                                                                                                                                                        23-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                              them, useful
                                                                    28-SEP-1998;
                                                                                      25-NOV-1998
                                                                                                            23-DEC-1998
                                                                                                                                      23-DEC-1998
                                            09-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nvention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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Matches
                                                                                                                                                                                                                                                                                                                                         New
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diagnostic primers and probes. Sequences AAA23463.A23502 represent hybridisation probes which may be used to isolate the cDNA clones of the
                                                                                                                                                                                   Gaps
                                                                                                                                                                                     ö
                                                                                                                                 1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2e+02; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                  945 GTGTGAGCGCAGACTGC 961
                                                                                                                                                                                                                                                          Greadacacacacacacac
                                                                                                                                                                                                                                                                                                                                                                   AAZ94157 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JUN-2000 (first entry)
                                                                                                                                                   Local Similarity 88.2
les 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                              AAZ94157;
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Phosphatidylethanolamine N-methyltransferase-2; PEWT2; human; liver cancer; hepatoma; antitumour; antiproliferative; therapy; diagnosis; PCR primer; ss.

Human PEMT2 PCR primer.

RESULT 293

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AAZ94157/

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The present sequence is that of a primer used in the PCR amplification of the open reading frame of a cDNA clone (see AAZ94150) encoding human phosphatidylethanolamine N-methyltransferase-2 (PEMT-2, see AAY79199). The PCR product was subcloned into mammalian expression vector pCI, and PEMT-2 was expressed in rat hepatona McArdle-RH777 cells. The invention relates to novel human PEMT2 polynucleotides and protein (see AAY79199), and to methods of using them in the treatment and diagnosis of liver disorders, such as liver cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polyribonucleotide solid phase synthesis; diagnosis; hybridisation; protein production modulation; 2'-deoxyfuranosyl moiety; anti-HIV; antiarteriosclerotic; nuclease resistant; atherosclerosis; AIDS; abnormal cell proliferation; tumour formation; ss.
                                                                                                                                                                                                       Isolated nucleic acid molecule encoding phosphatidylethanolamine N-methyltransferase protein used to treat phosphatidylethanolamine N-methyltransferase-associated disorders such as liver cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2e+02; ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 2 A; 7 C; 7 G; 3 T; 0 other;
                                                                                                                             (RESE ) RESEARCH CORP TECHNOLOGIES INC.
                                                                                                                                                                                                                                                            Example 8; Page 57; 111pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GTGGCCCAGCAGCAGGG 731
                                                                                                                                                          Cui Z;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ48149 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oligonucleotide SEQ ID NO:33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95US-0468037.
90US-0463358.
90US-0566977.
91WO-US05720.
92US-0835932.
                                                                                                      98US-0146218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GTAGCCCAGCAGCCGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98US-0035357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                       Walkey CJ,
                                                                                                                                                                                WPI; 2000-256956/22.
                                                                                                      02-SEP-1998;
                                                                           13-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-MAR-2000
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05-MAR-1992;
01-JUL-1992;
                                                 16-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US6005087-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                       Vance DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ48149;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 294
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à
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(ISIS-) ISIS PHARM INC.

Cook PD;

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14-JUN-2001
                                Best Loca
Matches
                                      RESULT 295
                                       AAS11124/c
ΩP
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The present invention describes nuclease resistant oligonucleotides (I) comprise comprising 2.* fluoro modified ribofuranosy1 nucleotides. (I) comprise covalently bound nucleotides, where the nucleotides are joined together by: (a) internucleotide linkages such that the base portion of the nucleotides forms a mixed base sequence; and (b) at least one of the nucleotides includes a modified ribofuranosy1 group bearing a 2.* fluoro substituent; provided that at least two of the nucleotides are 2.* fluoro substituent; provided that at least two of the nucleotides are 2.* fluoro substituent; provided that at least two of the nucleotides are 2.* fluoro compliced ribofuranosy1 nucleotides when the internucleotide linkages are phosphodiester nucleotides. (I) bind to their target mRNA and inhibit its expression. (I) are resistant to nuclease degradation and hybridise with the appropriate strength and fidelity to its target RNA/DNA. (I) are also useful as research agents, diagnostic agents and as oligonucleotide therapeutics. (I) may be used to treat atherosclerosis following the ungloblasty to prevent reocclusion of the treated arteries. (I) may also be used in conjunction with AZT to treat AIDS patients. (I) have been cused to modulate the expression of RAF gene, a cellular gene whose the conjunction implicated in abnormal cell province and a conjunction and province in a phonomal cell province of the invarience.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tumour formation. (1) are also used to modulate the expression of protein kinase C. (1) exhibit hybridisation properties of higher quality than phosphorous modified oligonucleotide duplexes containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          methylphosphonates, phosphoramidates and phosphate triesters. The present sequence represent an oligonucleotide used in the exemplification of the
Nuclease resistant oligonucleotides useful as research agents, diagnostic agents, and in the treatment of atherosclerosis and AIDS
                                                                                                                                                                                                       Example 19; Column 40; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present invention.
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0
                                                                       Gaps
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                                1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2e+02; cive 0; Mismatches 2; Indels
Sequence 19 BP; 13 A; 4 C; 2 G; 0 U; 0 other;
                                                                   Conservative
                                                 Local Similarity
                                                                 15;
                                Query Match
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1141 GCCTTTTTTTTTTTG 1157
                     ო
                  19
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AAS11124 standard; DNA; 19 (first entry) 24-OCT-2001 AAS11124;

Bacterial 16s RNA antisense oligomer #90.

Antisense; bacterial 16s ribosomal RNA; rRNA; bacterial infection; human; food grain supplement; livestock; poultry; therapeutic; ss.

Enterococcus faecium

29-NOV-2000; 2000WO-US42391.

99US-0168150. 29-NOV-1999;

AVIB-) AVI BIOPHARMA INC

WPI; 2000-072074/06. Kawasaki AM,

ВЪ.

WO200142457-A2.

[versen PL;

ô ribosomal RNA (rRNA) antisense oligomers. These sequences are antibacterial compounds comprising substantially uncharged antisense oligomers containing 8-40 nucleotide subunits, including a targeting nucleic acid sequence at least 10 nucleotides in length which is complementary to a bacterial 16 sor 23S rRNA nucleic acid sequence. The antisense oligomers are used for treating a bacterial infection in a human or a mammalian animal produced by Escherichia coli, Salmonella typhimurium, Pseudomonas aeruginosa, Vibrio cholera, Neisseria choorsthoea, Helicobacter pylori, Bartonella henselae, Haemophilus influenza, Shigella dysenterae, Staphyloococus aureus, Mycobacterium tuberculosis, Streptococcus pneumoniae, Treponema palladium and Chlamydia trachomatis. The antibacterial compound may be used as a food grain supplement in livestock and poultry food composition. Antibacterial compound, useful for treating bacterial infections and as livestock and poultry food supplement, comprises antisense oligonucleotides complementary to bacterial 16S and 23S rRNA -Gaps AAS11035-AAS11157 represent the coding sequences of bacterial 16S ·; 1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2e+02; ive 0; Mismatches 2; Indels Sequence 19 BP; 4 A; 5 C; 8 G; 2 T; 0 other; Disclosure; Page 35; 62pp; English 685 TTTGGGAGCCAGCGGC 701 Conservative WPI; 2001-457295/49. Local Similarity 15; Query Match Best Loca Matches ð

17 rrrregagecagecece g

AAH59922 standard; DNA; 19 AAH59922; AAH59922 

BP.

(first entry) 10-SEP-2001

Cyclin F ribozyme binding site SEQ ID NO:2346.

Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antieborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; virucide; actopic dermatities; actinic keratosis; squamous cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; se.

Homo sapiens Synthetic

WO200130362-A2

03-MAY-2001.

26-OCT-2000; 2000WO-US29500.

99US-0161532 26-OCT-1999;

(IMMU-) IMMUSOL INC.

Tritz R;

Robbins JM,

WPI; 2001-300427/31.

Treating proliferative skin or eye diseases and scarring, using

us09904568-3.rng

White RL;

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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (WMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, and expending (I). (I) can have antipsoriatic, and experimentation optimal mological, cytostatic, antiseborrheic, antidiabetic, antidiation, cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn car. AMHS7577 to AMH62099 represent sequences used in the exemplification of the present invention.
ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hedge PJ, Joslyn G;
s A, Vogelstein B;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Adenomatous polyposis coli; APC; human; neoplastic tissue; mutation detection; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single stranded conformational polymorphism; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Groden J, Hedg
, Thliveris A,
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 3 A; 6 C; 6 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DP1, SRP19, DP25 gene SSCP primer #11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Carlson M, G
Nakamura Y,
                                                               Example 1; Page 242; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             877 GCCAAGITCCAGGAGCT 893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABS67159 standard; DNA; 19 BP.
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91GB-0000963.
91GB-0000974.
91GB-0000975.
91US-0741940.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 GCCAGCTTCCAGGAGCT
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Markham AF,
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08-AUG-1991;
12-AUG-1994;
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Local S...
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16-JAN-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Albertsen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABS67159;
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The present invention relates to a method for haplotyping the human phospholipid transfer protein (PLTP) gene, involving determining the identity of the nucleotide present at one or more of the 25 polymorphic sites within the gene. This can be used to aid drug development for the
                                                                                                                                                                                      This invention relates to a novel method to aid in the diagnosis or prognosis of a neoplastic tissue of a human. The method involves detecting sometic alteration of wild-type adenomatous polyposis coli) protein in a tumour tissue isolated from a human (the alteration indicating neoplasia of the tissue). The method of the invention is useful in diagnosis or prognosis of a neoplastic tissue of a human. the method is useful in detection of genetic predisposition to cancer. The present sequence represents a DNA sequence used in the method
                                                                Method to aid in the diagnosis/prognosis of neoplastic tissues in humans, by detecting somatic alteration of wild-type APC protein in tumor tissue isolated from human, the alteration indicating neoplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, phospholipid transfer protein, PLTP; SNP, atherosclerosis,
single nucleotide polymorphism, high-density lipoprotein metabolism,
allele-specific oligonucleotide, probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human phospholipid transfer protein gene ASO probe SEQ ID NO: 20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genotyping phospholipid transfer protein gene of individual for haplotyping individual's gene, comprises determining identity of nucleotide pair at polymorphic sites for two copies of PLTP gene
                                                                                                                                                                                                                                                                                                                                                                                                                                              .
                                                                                                                                                                                                                                                                                                                                                                                                       1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 0 A; 4 C; 8 G; 7 T; 0 other;
                                                                                                                                                          Example 15; Column 31-32; 140pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100 ACAACCCCGGAGGCGCA 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-MAR-2000; 2000US-192127P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17 ACAACCCAGGAGCCGCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABA81571 standard; DNA; 15
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present in the individual
                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                  WPI; 2002-641559/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-662922/76.
                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
les 15; Conserv
                                                                                                                                                                                                                                                                                                                                   the invention
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                                                                                                                         of the tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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Matches
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AAS94583 standard; DNA; 15 BP.

AAS94583

(first entry)

14-FEB-2002

AAS94583;

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                 gene, possibly including atherosclerosis. The present sequence is an allele-specific probe used for detecting polymorphisms in the PLTP gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; ss; allele specific oligonucleotide; primer;
single nucleotide polymorphism; SNP; lipase endothelial isogene; LIPG;
drug screening; atherosclerosis; cardiovascular disorder;
LIPG haplotyping; LIPG genotyping.
 treatment of diseases associated with different haplotypes of the PLTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel genetic variants of Lipase, Endothelial isogenes, useful for improving efficiency and reliability in drug development for treating diseases associated with LIPG activity, e.g. atherosclerosis
                                                                                                                                                 Gaps
                                                                                                                                                 0
                                                                                                          Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                     Human LIPG gene allele specific oligonucleotide primer 39,
                                                                                                                                               0; Indels
                                                                                                         Query Match 1.0%; Score 13.6; DB 1; Best Local Similarity 92.9%; Pred. No. 1.6e+02; Matches 13; Conservative 1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 BP; 1 A; 4 C; 7 G; 2 T; 1 other;
                                                                      Sequence 15 BP; 6 A; 2 C; 5 G; 1 T; 1 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 16; Page 14; 134pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kazemi A, Kliem SE,
                                                                                                                                                                                                                                                                                                           ABL91860 standard; DNA; 15 BP.
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                                                                                                                                                                                 918 AAAGGAGATGGCAG 931
                                                                                                                                                                                                      2 AAAGGARATGGCAG 15
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8 x 6 6 6 6
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The invention relates to single nucleotide polymorphisms in the gene encoding the human phospholipid transfer protein (PLTP). A method for haplotyping the PLTP gene in an individual comprises identifying the nucleotide at one or more polymorphic sites and determining whether one of the specification or whether both copies are defined by a haplotype given in the specification or whether both copies are defined by a haplotype pair. This method is useful in genotyping, whereby all possible haplotype pair. This method is useful in genotyping, whereby all possible haplotype pair and a haplotype or haplotype pair of the haplotype or haplotype pair in a reference population, where a higher haplotype or haplotype pair in a reference population, where a higher haplotype or haplotype for haplotype or haplotype for nation indicates the trait is associated with the haplotype or haplotype pair. Purp and its corresponding DNA are used for studying the expression and function of PUTP, for use in screening for candidate drugs to treat diseases related to PLTP activity. The sequences are also useful for studying the effect of variation on the biological activity of PLTP as well as on the binding affinity of candidate drugs targeting PLTP for treating atherosclerosis. Sequences ASS94691 represent allele-specific oligomucleotide probes, sequencing primers and PCR primers used for detecting PLTP gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
0
                                                                                                                                                           Human, phospholipid transfer protein, PLTP; haplotyping, haplotype pair;
single nucleotide polymorphism; genotyping; gene therapy, drug screening;
binding affinity; atherosclerosis; ss; sequencing primer; PCR primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated polynucleotide which is polymorphic variant of phospholipid transfer protein (PLTP) gene, having any one of polymorphic sites PSI-PS25, for studying function of PLTP, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .
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                                                                                                                      Human PLTP gene allele-specific oligonucleotide probe #17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 6 A; 2 C; 5 G; 1 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 15; Page 70; 99pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                         26-MAR-2001; 2001WO-US09776.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expressing PLTP protein
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                                                                                                                                                                                                                                                                                                         WO200172966-A2.
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                                                                                                                                                                                                                              probe.
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RESULT 301 AAC67429

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Gaps ٥.

0; Indels

1.0%; Score 13.6; DB 1; 92.9%; Pred. No. 1.6e+02; live 1; Mismatches 0;

534 GCAGCTGGGTGCCC 547 

ð g

13; Conservative

Matches

Local Similarity

Query Match

Length 15;

RESULT 300

WO9966075-A2

23-DEC-1999

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of or diagnosing Alzheimer's disease using single nuclectide polymorphisms (SNPs) present in an individual's mitochondrial DNA (mtDNA). In additon, the SNPs identified can be used to identify agents suitable for use in treating Alzheimer's disease. Sequences AAC67301-C67610 are PCR primers used to demonstrate the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present invention describes a novel method for determining the risk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; mitochondrial DNA; extramitochondrial DNA; mtDNA; exmtDNA; diagnosis; quantification; detection; dystonia; Alzheimer's disease; Huntington's disease; barkinson's disease; schizophrenia; stroke; non-insulin dependent diabetes mellitus; mitochondrial encephalopathy; lactic acidosis; myoclonic epilepsy ragged red fibre syndrome; Leber's hereditary optic neuropathy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Diagnosing a subject at the risk for or having Alzheimer's disease comprises determining at least one single nucleotide polymorphism in mitochondrial DNA associated with the disease in the sample from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human mitochondrial DNA NADH dehydrogenase PCR primer SEQ ID NO:76.
                                                                                                                                                                                                                                                       Human; mitochondrial genome; single nucleotide polymorphism; SNP; Alzheimer's disease; mtDNA; PCR primer; ss.
                                                                                                                                                                                    Alzheimer's disease-linked mitochondrial SNP PCR primer #129.
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80.0%; Pred. No. 2.5e+02;
ative 0; Mismatches 4;
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                                                                                                                              (first entry)
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AAC67429 standard;
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Best Local Similarity
Matches 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-APR-1999;
22-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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The present invention describes a method for the quantification of extramitochondrial DNA (exmtDNA) by determining the ratio of a first and search biological sample containing exmtDNA and mitochondrial DNA (m.DNA) to determine the risk or presence of a disease associated with altered mitochondrial function. The method can be used to determine the risk of or presence of a disease associated with altered mitochondrial function, especially Alzheimer's disease, Huntington's disease, Parkinson's disease, dystonia, schizophrenia, non-insulin dependent diabetes mellitus, mitochondrial encephalopathy, lactic dependent diabetes mellitus, mitochondrial encephalopathy, lactic acidosis, stroke, myoclonic epilepsy ragged red fibre syndrome and leber's hereditary optic neuropathy. The method can also be used to identify agents sultable for treating such diseases, in particular Alzheimer's disease. AAZ57202 to AAZ57313 represent nucleotide sequences used in the exemplification of the present invention. More specifically and such and such also be used to season in the exemplification of the present invention. More specifically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RT-PCR primer; DNA sequence determination; gene sequence analysis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ57206 to AAZ57313 are PCR primers used in the detection of exmtDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                          Quantification of extramitochondrial DNA for diagnosis of, e.g. Alzheimer's, Huntington's and Parkinson's disease
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                                                                                                                                                                                                    Davis RE;
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                                                                                                                                                                                                  Fahy ED,
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                                                                                                                                                                                                Clevenger W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RT-PCR primer of the invention SEQ ID 5.
                                                                                                                                                                                                                                                                                                            Disclosure; Page 32; 157pp; English
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98US-0098079.
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                                                                                                                                                                                              Ghosh SS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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es 16; Conserv
                                                                                                                                                              (MITO-) MITOKOR,
                                                                                                                                                                                                Herrnstadt C,
                                                                14-JUN-1999;
                                                                                             L5-JUN-1998;
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Length 21; 4; Indels

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us09904568-3.rng

Page 148

Peptides having at least two new nucleotides - useful as primers in Disclosure; Page 10; 19pp; Japanese. WPI; 1999-183822/16

This sequence represents a primer of the invention. The invention relates to sequences of at least two nucleotides of formula:

(X)m5'-(alphain-beta-N3'; or (X)m5'-(gamma)k-delta-N3'; where x = alabelled compound and/or a nucleotide with voluntary sequence; m = 0 or 1; alpha = thymine; n = natural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, guanine or cytosine; N = adenine, guanine, cytosine or thymine; gamma = thymine; k = natural number of 3 or over indicating the repetition of gamma, in which thymine expressed by gamma is composed of 1/3 or less of adenine, guanine and/or cytosine. The new nucleotides are useful as primers for XT-PCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene sequences.

Sequence 15 BP; 0 A; 0 C; 2 G; 13 T; 0 other;

1.0%; Score 13.4; DB 1; 93.3%; Pred. No. 1.8e+02; tive 0; Mismatches 1; 1144 TTTTTTTTTTGG 1158 1 TTTTTTTTTTGG 15 Local Similarity 93.3 Query Match Matches à g

AAF95031 standard; DNA; 15 RESULT 304 AAF9503

AAF95031;

23-MAY-2001

Mutant capture oligonucleotide #24.

Tubercle bacillus; drug sensitivity; drug resistance; rifampicin; streptomycin; kanamycin; isoniazid; ethambutol; rpoB gene; rrs gene; rpst gene; inhA gene; katG gene; embB gene; probe; PCR primer; ss.

Mycobacterium tuberculosis.

EP1076099-A2

14-FEB-2001.

02-AUG-2000; 2000EP-0306563

99JP-0220357 03-AUG-1999;

(NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.

Takenishi S; Suzuki Y, Nishida M,

WPI; 2001-246696/26.

New oligonucleotides, nucleic acid probes and primers are useful for differentiating drug-resistance and determining infection with tubercle bacilli

Claim 10; Page 25; 114pp; English.

The present invention relates to oligonucleotides based on nucleotide sequences obtained from both wild-type tubercle bacilli (wtTB) that are susceptible to a drug and mutant-type tubercle bacilli (mtTB) that are resistant to a drug. The drugs used in the present invention are rifampicin (RFP), streptomycin (SM), kanamycin (KM), isoniazid (INH) and

ethambutol (EB). The rpoB gene is responsible for resistance to RFP; the responsible for resistance to SM and KM; the rpsL gene is responsible for resistance to SM; the inhA gene is responsible for resistance to INH, the katG gene is responsible for and the embB gene is responsible for resistance to INH; and the embB gene is responsible for resistance to EB. The present invention also relates to nucleic acid probes having part of a nucleotide sequence of tubercle bacilli (TB) responsible for drug resistance and primers used to generate the probes. The present sequence is an present invention can be used to enable the differentiation of drug resistance and the determination of infection with tubercle bacilli simultaneously.

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Sequence 15 BP; 3 A; 5 C; 5 G; 2 T; 0 other;

Gaps . 0 Length 15; 1; Indels Query Match
1.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 1.8e+02;
Matches 14; Conservative 0; Mismatches 1;

0

à

RESULT 305 AAF45161

0

Gaps

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Length 15;

AAF45161 standard; RNA; 15 BP

30-MAR-2001 (first entry)

Antisense oligonucleotide #10.

Antisense therapy, antiproliferative, antinflammatory, antipsoriatic; cytostatic; dermatological, cardiant, virucide; ophthalmological, keloid, skin disorder, Insulin-like Growth Factor. I receptor; IGF-1; pityriasis; IGF binding procein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hypermeovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.

Homo sapiens.

WO200078341-A1

21-JUN-2000; 2000WO-AU00693.

99US-0140345. 21-JUN-1999;

(MURD-) MURDOCH CHILDRENS RES INST.

Edmondson SR; Werther GA, Wraight CJ,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation

Claim 15; Page 115; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBB], which is capable of inhibiting or reducing growth factor maddated cell proliferation, inflammation and/or other disorders. The present sequence is one such antisense oligonuclectide. The method is useful for ameliorating the

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effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia.
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Sequence 15 BP; 4 A; 5 C; 6 G; 0 U; 0 other;

1.0%; Score 13.4; DB 1; Length 15; 93.3%; Pred. No. 1.8a+02; Ive 0; Mismatches 1; Indels 552 GGCAGGCATGCACAC 566 93.38; Conservative Local Similarity Les 14; Conserv Query Match Matches ð

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Gaps

.; 0

1 GGCAGGCAGCACAC 15 g

AAF46436

AAF46436 standard; DNA; 15 BP

AAF46436;

IGFBP2 oligonucleotide #1275.

30-MAR-2001 (first entry)

Antisense therapy, antiproliferative, antinflammatory, antipsoriatic, cytostatic, dermatological, cardiant, virucide, ophthalmological, keloid, skin disorder, Insulin-like Growth Factor I receptor, IGF-1, pityriasis, IGF binding procein, IGFBP-2, IGFBP3, inflammation, psoriasis, pilaris; growth factor mediated cell proliferation, ichthyosis; serborrhoea, ruba, keratosis, neoplasia; scleroderma; wart, skin cancer, sclerotic disease, hypermeovascular condition, hyperplasia, kidney disease, neovascular condition of the retina; ss.

WO200078341-A1.

28-DEC-2000

21-JUN-2000; 2000WO-AU00693.

99US-0140345. 21-JUN-1999;

(MURD-) MURDOCH CHILDRENS RES INST.

Werther GA, Edmondson SR; Wraight CJ,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation

Example 6; Page 42; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [167]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonuclectide which can be used to design the antisense oligonuclectides of the present invention (see AAF45151 and AAF45153-F55161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, plaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition as a meovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other 

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                                    Gaps
sclerotic disease, kidney disease, hyperproliferation of the inside blood vessels or any other hyperplasia.
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                           Length 15;
                    ; DB 1; her,
1.8e+02;
1; Indels
              Sequence 15 BP; 3 A; 5 C; 4 G; 3 T; 0 other;
                         1.0%; Score 13.4; I 93.3%; Pred. No. 1.8e
                                                                                 ВР
                                                                                                               IGFBP2 oligonucleotide #1276.
                                              253
                                                       Н
                                                                               AAF46437 standard; DNA; 15
                                                                                                     (first entry)
                                    Conservative
                                             239 CATCTGCATCTGGGA
                                                        15 CATCTGCAGCTGGGA
                              Similarity
                       Query Match
Best Local Simil
Matches 14; C
                                                                                                     30-MAR-2001
                                                                                          AAF46437;
                                                                       RESULT 307
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Antisense therapy, antiproliferative, antiinflammatory, antipsoriatic, cytostatic, dermatological; cardiant; virucide, ophthalmological; keloid, skin disorder; Insulin-like Growth Factor. I receptor; IGF-1; pitchriaais; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hypermeovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.

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WO200078341-A1.

28-DEC-2000,

21-JUN-2000; 2000WO-AU00693

21-JUN-1999;

(MURD-) MURDOCH CHILDRENS RES INST.

99US-0140345,

Edmondson SR; Werther GA, Wraight CJ,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation

Example 6; Page 42; 201pp; English.

The present invention relates to a method for ameliorating the effects
of skin disorders. The method comprises contacting the skin with an
antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1
receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of
inhibiting or reducing prowth factor mediated cell proliferation,
inflammation and/or other disorders. The present sequence is an
oligonuclectide which can be used to design the antisense
oligonuclectide which can be used to design the antisense
oligonuclectide which can be used to a mealiorating the effects of
psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids,
keratosis, neoplasis, pityriasis, ruba, pilaris, serborrhoea, keloids,
keratosis, neoplasis, cleroderma, warts, benign growths, cancers of the
skin, a hyperneovascular condition such as a neovascular condition of the
retina, brain or skin, growth factor-mediated malignancies, other
condition of the inside of blood vessels or any other hyperplasia.

Sequence 15 BP; 3 A; 6 C; 4 G; 2 T; 0 other;

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                                                                                                                                                                                                                                                                                                                                                            Antisense therapy, antiproliferative, antinflammatory, antipsoriatic, cytostatic, dermatological, cardiant, virucide, ophthalmological, keloid, skin disorder, Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein, IGFBP-2; IGFBP3; inflammation, psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba, keratosis, neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hypermeovascular condition; hyperplasia; kidney disease; neoblation of the recina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell
                                                     Gaps
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                     Length 15;
                                                   Indels
                 Score 13.4; DB 1;
Pred. No. 1.8e+02;
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                                                 0; Mismatches
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             1.0%;
                                                                                      238 GCATCTGCATCTGGG 252
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                                                                                                                        15 GCATCTGCAGCTGGG 1
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The 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                            Antisense therapy, antiproliferative, antinflammatory, antipsoriatic, cytostatic, dermatological, cardiant, virucide, ophthalmological, keloid, skin disorder, Insulin-like Growth Factor I receptor; IGF-1, pityliasis; IGF binding proctein, IGFBP-2, IGFBP3, inflammation, psoriasis; pilaris, growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba, keratosis; neoplasia; scleroderma; wart, skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia, kidney disease; necovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oligonacleotides of the present invention (see AAF45151 and AAF45153 F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serbornhoea, keloids, keratosis, neoplasias, soleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the sclero, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kindy disease, hyperproliferation of the inside of blood vessels or any other hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligomucleotide. (for Insulin-like Growth Factor [IGF] -1 receptor, IGF binding protein [IGFBP] -2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligomucleotide which can be used to design the antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 1.0%; Score 13.4; DB 1; Length 15; Local Similarity 93.3%; Pred. No. 1.8e+02; es 14; Conservative 0; Mismatches 1; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 3 A; 2 C; 8 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Edmondson SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MURD-) MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 6; Page 42; 201pp; English.
                                                                                                                                                                                                                                                               IGFBP2 oligonucleotide #1342.
  251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                888 GGAGCTGCGGTACAG 902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-JUN-2000; 2000WO-AU00693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GGAGCTGGGCTACAG 15
                                                                                                                                        AAF46503 standard; DNA; 15
                                                                                                                                                                                                                      (first entry)
                     15 GGCATCTGCAGCTGG
237 GGCATCTGCATCTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Werther GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-041421/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200078341-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-JUN-1999;
                                                                                                                                                                                                                      30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wraight CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-DEC-2000
                                                                                                                                                                              AAF46503;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Loca
Matches
                                                                                                                    AAF46503
                                                                                                                                                           g
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0

0; Gaps

1.0%; Score 13.4; DB 1; Length 15; 93.3%; Pred. No. 1.8e+02; ve 0; Mismatches 1; Indels

1 Similarity 93.3%; 14; Conservative

Best Local Similarity

Matches

Query Match

BP.

AAF49864 standard; DNA; 15

AAF49864;

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Antisense therapy, antiproliferative, antiinflammatory, antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin discorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 0 A; 5 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                                     neovascular condition of the retina; ss
                                                                                                                                                                                                                                                                                                                                                                                                    Wraight CJ, Werther GA, Edmondson SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; Page 66; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                           (MURD-) MURDOCH CHILDRENS RES INST,
                          AAF49863 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                       21-JUN-2000; 2000WO-AU00693.
                                                                                                     IGF-I oligonucleotide #823.
                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-041421/05.
                                                                                                                                                                                                                                                                       WO200078341-A1.
                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                 21-JUN-1999;
                                                                           30-MAR-2001
                                                                                                                                                                                                                                                                                                28-DEC-2000.
                                                   AAF49863;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
RESULT 310
             AAF49863/c
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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFB]-2 or IGFBB3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense and AAF451S1-The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, seforthoea, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia.
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                                Gaps
                                 .
    Length 15;
                              Indels
1.0%; Score 13.4; DB 1;
93.3%; Pred. No. 1.8e+02;
iive 0; Mismatches 1;
                                                   553 GCAGGCATGCACACA 567
                                                                             15 GCAGGCAGGCACACA 1
                             Conservative
            Similarity
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14;

Matches

à

RESULT 311 AAF49864/C

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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1.

receptor, IGF binding protein [IGFB]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonuclectide which can be used to design the antisense oligonuclectide which can be used to design the antisense oligonuclectide which can be used to design the effects of AAP45153-F45161. The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma warts, benign growths, cancers of the skin, a hypernevascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other celina, brain or skin, growth factor-mediated malignancies, other condition of the inside of blood vessels or any other hyperplasia.
                                                                                                                                                Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neoblation of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15 BP; 0 A; 6 C; 5 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Edmondson SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MURD-) MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 8; Page 66; 201pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  552 GGCAGGCATGCACAC 566
                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-JUN-2000; 2000WO-AU00693,
                                                                                                               IGF-I oligonucleotide #824.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF51703 standard; DNA; 15
                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Werther GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-041421/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                      WO200078341-A1.
                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wraight CJ,
                                                                          30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                             28-DEC-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 312
AAF51703/c
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Matches
EXXXX
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30-MAR-2001 (first entry)

IGF-I oligonucleotide #2663

Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pitzriasis; IGF binding protein; IGFBP-2; IGFBF3; inflammation; psoriasis; piratis; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scaleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.

Homo sapiens.

WO200078341-A1.

28-DEC-2000.

21-JUN-2000; 2000WO-AU00693

99US-0140345 21-JUN-1999; (MURD-) MURDOCH CHILDRENS RES INST.

Edmondson SR; Wraight CJ, Werther GA,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 8; Page 78; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGR]-1 receptor, IGR binding protein [IGRBP]-2 or IGRBP3, which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AF4545152 Pf5161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis; ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, Sclaroderma, warts, benign growths, cancers of the seting and a provise of the seting and a retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia. 

Sequence 15 BP; 4 A; 2 C; 6 G; 3 T; 0 other;

Gaps ; 0 1.0%; Score 13.4; DB 1; Length 15; 33.3%; Pred. No. 1.8e+02; ve 0; Mismatches 1; Indels Indels L Similarity 93.3%;
14; Conservative Sest Local Similarity Query Match **Jatches** 

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CGCGCTGATCCTCAA 172

Crcccrcarcara 1 15

à Пb

ABS97176 standard; DNA; 15 ABS97176; RESULT 313 ABS97176,

BP.

(first entry)

23-DEC-2002 XXXXXXXX

Human CYP4501A2 Exon 4 and 5 sequencing primer #4.

The state of the s primer; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;

Homo sapiens.

WO200257410-A2.

25-JUL-2002.

28-NOV-2001; 2001WO-US44838.

28-NOV-2000; 2000US-0724389.

(DNAS-) DNA SCI LAB INC.

Guida M, Hall J;

WPI; 2002-698522/75.

Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related traits

Example 2; Page 101; 714pp; English.

This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP450121), adrenergic receptor beta1 (PABRI), cytochrome P450 O21 (CYP450121), adrenergic receptor beta1 (ADBRI), cytochrome P450 O21 (CYP450121), adrenergic receptor beta1 (ADBRI), cytochrome P450 O21 (CYP450121), adrenergic receptor nuclear translocator (ARN), cathepsin S (CTSS), cyclooxgenase 2 (CXCZ), diazepam binding inhibitor (DBI), epoxide hydrocarbon receptor nuclear translocator (ARNT), cathepsin S (CTSS), cyclooxgenase 2 (CXCZ), diazepam binding inhibitor (DBI), epoxide hydrocylase 2 (EDRX2), 5-lipoxygenase 2 (ARNT), cathepsin S (CTSS), dutathione-S-transferase 12 (GYCTS), diazepam binding cativating protein (FARNT), NADPH quinone oxidoreducase 2 (NQC2), alstanie-N-methyl transferase 2BH (UGT2BH), UDP-glucuronosyl transferase 2BH (UGT2BH), UGT2BH, UDP-glucuronosyl transferase 2BH (UGT2BH), UDP-glucuronosyl transferase 2BH (UGT2BH), UDP-glucuronosyl transferase 2BH (UGT2BH), UGT2BH, UDP-glucuronosyl transferase 2BH (UGT2BH), UDP-glucuronosyl transferase 2BH (UGT2BH), UDP-glucuronosyl transferase 2BH (UGT2BH), UGT2BH, UDP-glucuronosyl transferase 2BH (UGT2BH), UDP-glucuronosyl transferase sontained individuals for altered secential individuals for altered secential individuals for altered secential individuals

prostate/colon tumour suppressor gene; allelic loss; colorectal cancers; microsatellite analysis; sequence tagged site; primer; probe; PCR; amplification; chromosome; ss.

Chromosomal locus E15 primer #2.

(first entry)

04-JUL-1996

BP.

AAT06918 standard; DNA; 17

New DNA encoding a prostate tumour suppressor protein - from chromosome 8, for the diagnosis and treatment of prostatic and colorectal cancer

94US-0246604

22-MAY-1995; 20-MAY-1994;

30-NOV-1995.

WO9532214-A1

Synthetic.

WB;

Isaacs

(CANJ-) CANJI Bookstein R, WPI; 1996-020526/02.

Disclosure, Page 86, 122pp; English.

us09904568-3.rng

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haematological function, in KLK2 for altered serine protease activity the prostate, in LTF for altered immunological or haematological function, in CHMR3, CHMR4 or CHMR5 for altered central and peripheral nervous system function. The present sequence represents a sequencing primer used to sequence the polymorphic genes of the invention.
                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                            Human, neuropeptide Y; NPY, isogene; SNP; atherosclerosis; obesity; psychological disorder; single nucleotide polymorphism; alcoholism; antiarteriosclerotic; anorectic; PCR; primer; ss.
                                                                                                                             ö
                                                                                                 Query Match 1.0%; Score 13.4; DB 1; Length 15; Best Local Similarity 93.3%; Pred. No. 1.8e+02; Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                   Human neuropeptide Y allele specific primer SEQ ID NO: 50.
                                                                          Sequence 15 BP; 3 A; 2 C; 9 G; 1 T; 0 other;
                                                                                                                                                      582 CCTCCGTCTGCCCCC 596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-DEC-2000; 2000WO-US34758.
                                                                                                                                                                             cercagrerecede
                                                                                                                                                                                                                                           AAL48126 standard; DNA; 15
                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                        WO200251857-A1.
                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-JUL-2002.
                                                                                                                                                                                                                                                                                              27-SEP-2002
                                                                                                                                                                              15
                                                                                                                                                                                                                                                                    AAL48126;
                                                                                                                                                                                                                   RESULT 314
                                                                                                                                                                                                                                AAL48126
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primers AAT06887-932 were used to analyse the breakpoints at chromosomal locus 8p22-21, contained in patients having prostate cancer, by microsatellite analysis and sequence taggaded sites (STS). The region contains a prostate/colon tunour suppressor gene (PTSG). The primers and amplified fragments were used to screen a YAC library of prostate diagnosis and treatment of prostate and colorectal cancer by the pigg (AAT06880), which can be used in the diagnosis and treatment of prostate and colorectal cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Prostate/colon tumour suppressor; allelic loss; prostate cancer; colorectal cancer; microsatellite analysis; sequence tagged site; STS; amplification; chromosomal location 8q22-21; probe; primer; gene mapping; diagnosis; treatment; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer E15 for mapping prostate/colon tumour suppressor gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 93.3%; Pred. No. 2.1e+02;
14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 1 A; 4 C; 6 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT64712 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 237 GGCATCTGCATCTGG 251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 GGCATCTGCTTCTGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Simil
Matches 14; C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
12-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT64712;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT64712
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Synthetic. Homo sapiens.

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Gaps

. 0

Indels

1.0%; Score 13.4; DB 1; Length 15; 3.3%; Pred. No. 1.8e+02;

/ Match Local Similarity 93.3%; Pred. No. 1.00

Query Match

Matches

639 GCTCTGCATCCCCCA 653

ð

GCTCTGAATCCCCCA 15

RESULT 315 AAT06918

Sequence 15 BP; 3 A; 7 C; 2 G; 3 T; 0 other;

present sequence is NPY coding sequence

The present invention provides the human neuropeptide Y (NPY) gene and single nucleotide polymorphisms (SNPs) identified therein. The sequence can be used in the treatment of disorders associated with NPY, including atherosclerosis, obesity, psychological disorders and alcoholism. The present sequence is an allele specific primer used to isolate the human

New genetic variants of the human Neuropeptide Y (NPY) gene useful for treating disorders affected by abnormal expression or function of NPY isogene e.g., atherosclerosis or obesity  $\dot{\cdot}$ 

Claim 11; Page 17; 80pp; English.

S,

Stephens

Denton RR, Lanz EM, Nandabalan K,

WPI; 2002-566671/60.

Chew A,

(GENA-) GENAISSANCE PHARM INC.

21-DEC-2000; 2000WO-US34758.

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JP09098790-A

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Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiniflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss.
                                                                                                                                                                                                                                                                                                                                               The present primer was used in the mapping of a gene encoding 2 forms of a prostate/colon tumour suppressor (P/CTS). The P/CTS gene was isolated by analysis of allelic loss in patients with prostate cancer, and was putatively located to the chromosomal location #Q22-21 via microsatellite analysis and the use of sequence tagged sites (STS). Primers and probes derived from the gene can be used to screen lambda cDNA libraries for genes encoding P/CTS form 1 and 2. The P/CTS or its cDNA can be used in the diagnosis and treatment of prostate and colorectal cancers.

(Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                   New prostate/colon tumour suppressor gene - mapped to a locus on human chromosome \vartheta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.1e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Integrin alpha 6 subunit substrate sequence SEQ ID NO:4572.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 1 A; 4 C; 6 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                      Disclosure; Page 26; 45pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA21346 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             237 GGCATCTGCATCTGG 251
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                                                                                   96JP-0062144
                                                                                                                  95US-0445515
                                                                                                                                                  (CANJ-) CANJI INC.
(UYJO ) UNIV JOHNS HOPKINS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14; Conservative
                                                                                                                                                                                                    Bookstein R;
                                                                                                                                                                                                                                  WPI; 1997-275447/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                CANJ-) CANJI
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                                                                                                                 22-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-0CT-1999
                                                                                   22-FEB-1996;
                                                 15-APR-1997
                                                                                                                                                                                                     Isaacs WB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA21346;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 317
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Gaps

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The present invention describes enzymatic nucleic acid molecules with CC RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNY) gene, an integrin subunit betea 3 gene, an integrin subunit betea 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA1762 to AAA1762 represent their corresponding target sequences; AAA1762 to AAA1762 to AAA1762 to AAA1762 to AAA1763 to AAA1763 to AAA1915 to AAA1916 subunit, and AAA2335 to AAA2130 and AAA215 to AAA2168 represent their corresponding target sequences for integrin subunit beta 3, and AAA22476 to AAA23343 to AAA2342 represent their corresponding target sequences. The ribozymes of the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially AANT, integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are considered to treat cancer, diabetric retinopathy, age related compaction (ARW), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, corresponding target sequence conscileroms of tubercoms, myopic degeneration, psoriasis, struge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2, integrin subunit alpha-6, or integrin subunit beta-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                             Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors \,
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                                                                          Coeshott C, McSwiggen JA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.0%; Score 13.4; DB 1; 0.0%; Pred. No. 2.1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11; Mismatches
                                                                                                                                                                                                                                                             Claim 55; Page 202; 305pp; English.
                                                                             Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAC66363 standard; DNA; 17 BP.
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                        (RIBO-) RIBOZYME PHARM INC.
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                                                                             Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                 WPI; 1999-591315/50.
                                                                             Pavco PA,
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Protection; pathogen infection; vaccination; immunisation; policvirus; Bordetella pertussis; respiratory syncytial virus; Mycoplasma pneumoniae; meningococous; preumococus; preumococus; preumococus; preumococus; corynebacterium dipttheriae; olostridium tetani; hepatitis B virus; Chlamydia pneumoniae; Chlamydia trachomatis; Moraxella catarrhalis; PCR primer used to amplify B. pertussis Sl DNA.

(first entry)

22-FEB-2001

Bordetella pertussis. WO200064457-A1. PCR primer; 12-NOV-2000, 

21-APR-2000; 2000WO-US10954.

99WO-US06507 98US-0079678

24-MAR-1999; 27-MAR-1998;

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Disclosure; Page 58; 111pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Housman DE,
                                            Claim 37; Page 67; 164pp; English
                                                                                                                                                                                                                                                                                                               ВР.
                                                                                                                                            protein and interferon alpha.
                                                                                                                                                                                       1.0%;
                                                                                                                                                                                                                                   CICCGICIGCCCCCC 597
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                                                                                                                                                                                                                                                                                                              AAA36112 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                             14; Conservative
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                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                          26-JUL-2000
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                                                                                                                                                                                                                                    583
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                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                         RESULT 320
                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                      AAA3611
                                                                                                                                                                                                                                                                                                                ò
                                                                                                                                                                        comprises a live commensal oral organism genetically modified to express multiple immunogenic fragments of the pathogen. The composition has antibacterial and antiviral activity and acts as a vaccine. The composition has composition which is administered orally or intranasally, is used for prophylactically treating a host against infection by a pathogen such as prochylactically treating a host against infection by a pathogen such as preming occores, respiratory syncytial virus, poliovirus, Mycoplasma premingluenza, Corynebacterium diphtheriae, Clostridium tetani, hepatitis parainfluenza, Corynebacterium diphtheriae, Clostridium tetani, hepatitis or their combinations. The composition can also be used for chronic their combinations. The composition can also be used for chronic sequence represents a PCR primer used to amplify a Bordetella pertussis DNA sequence, which is used in an example illustrating the use of the
                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                            which express immunogenic fragments of mucosal pathogens, used as oral vaccines to treat host against Bordetella pertussis, poliovirus
                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                 oral commensal bacteria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ribozyme; erythropoietin; granulocyte colony stimulating factor;
                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                           1.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.1e+02; tive 0; Mismatches 1; Indels
                                                                                                 Composition having genetically modified live
                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 2 A; 9 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               McSwiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hammerhead ribozyme substrate #504.
                                                                                                                                                       Example 1; Page 25; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zwick M, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                       760 CGGTGGCGGGTGGAT 774
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               99US-0298135
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 93.3
Matches 14; Conservative
                                   (UYDA-) UNIV DALHOUSIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     interferon alpha; ss.
                                                         Halperin SA;
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                                                                             WPI; 2000-687261/67.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
               23-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-0CT-2000
                                                                                                                                     infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF02209;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF02209
                                                         Lee SF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 319
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A method has been developed for detecting the presence or absence of a single nucleotide polymorphism (SNP) allele in a genomic sample. The method comprises preparing a reduced complexity genome (RCG) from the genomic sample and analysing the RCG for the presence or absence of a SNP allele. The method can be used to characterise a tumour, to generate a genomic pattern for an individual genome or to generate a genomic classification code for a genome. The method can be used to assess whether a subject is at risk for developing a disease or to identify a set of SNP alleles associated with a disease. The method can also be used to perform linkage analysis. AAA35944 to AAA35947 represent
                                                                                                                                                                                                                                                                                              The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRP-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; single nucleotide polymorphism; SNP; genotyping; DNA analysis; allele specific oligonucleotide; ASO; reduced complexity genome; RCG; genomic classification; identification; by fingerprinting; tumour characterisation; hybridisation; BS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human genomic SNP allele specific oligonucleotide SEQ ID NO:169.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.4; DB 1;
Pred. No. 2.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Charest A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 1 A; 10 C; 3 G; 3 T; 0 other;
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The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(di)thioate link, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A) that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or canderium), in vivo or by transforming cells ex vivo and implanting treated cells, or for other conditions associated with levels of oestrogen receptor. Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that creativition andonucleases are used with DNA). The combination of modifications in (A) improves residence to nucleases, binding affinity and/or activity. AAA23503 to AAA4747 represent oestrogen receptor carriesponding target sequences, and AAA26107 to AAA2618 represent certoper receptor hairpin ribozyme sequences, and AAA26107 to AAA2618 represent centroper their corresponding target sequences, AAA25619 to AAA2611 represent corresponding corresponding target sequences.
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sequences used in the exemplification of the present invention. AAA35948 to AAA36632 represent nucleotide sequences containing SNPs.
                                                                                                                                                                                                                                                                                                                                                                                                       Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:2072
                                                                                                                                                                                                                                                                                                                                                                                                                                          Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; hammedification; cancer; phosphorothioate; endonuclease; anticancer; breat cancer; endometrium cancer; ss.
                                                                                                                               Gaps
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                                                                                          Length 17;
                                                                                                                               1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Karpeisky A,
Haeberli P;
                                                                                          1.0%; Score 13.4; DB 1; 93.3%; Pred. No. 2.1e+02;
                                                       Sequence 17 BP; 2 A; 0 C; 2 G; 13 T; 0 other;
                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              McSwiggen JA, is T, Woolf T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 77; Page 83; 148pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Beigelman L, McSwig
Zwick M, Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequences, used to treat cancer
                                                                                                                                                                1145 TTTTTTTTTGGA 1159
                                                                                                                                                                                                                                                                                               AAA25574 standard; DNA; 17 BP.
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98US-0103636.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                             14; Conservative
                                                                                      Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                    19-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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23-JUN-1998;
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other ribozyme sequences and antisense oligonucleotides used in the exemplification of the present invention.

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The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCAL, BRA2, CFTR, cyclin-dependent Kinase inhibitor 2A (CDKMZA), APC, Factor V, Factor VIII, Factor IX, haemoglobin alpha locus (HBA1), haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6, apolipoprotein E (APC), presenilin-1 (PSBN1) and presenilin-2 (PSBN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, such as cancer, adenosine deaminase deficiency, cystic fibrosis, altahmer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting
                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                          Human; gene therapy, adenosine deaminase deficiency; p53; beta-globin, retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factox V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factox VII; Factox IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UGTI; syndrome; APP; PSEN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentlin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical
                                                                             Gaps
                                                                             .,
                                                                                                                                                                                                                                                                                                                                                                                     Adenosine deaminase deficiency correcting oligo SEQ ID NO: 35.
                                    Length 17;
                                                                             Indels
                                ch 1.0%; Score 13.4; DB 1; 1 Similarity 93.3%; Pred. No. 2.1e+02; 14; Conservative 0; Mismatches 1;
Sequence 17 BP; 6 A; 4 C; 1 G; 6 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rice MC;
                                                                                                                                                                                                                                                               ABA77189 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-MAR-2000; 2000US-192176P.
27-MAR-2000; 2000US-192179P.
01-JUN-2000; 2000US-208538P.
                                                                                                                   71 CATAGGATGAATAAT 85
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                                                                                                                                                      17 CATTGGATGAATAAT 3
                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antilipemic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200173002-A2.
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modification -
                                                                                                                                                                                                                                                                                                                                              24-JAN-2002
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Gaps ·.

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the invention.

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The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFRF, cyclin-dependent kinase inhibitor 2A (CDKNZA), APC, Factor VII, Factor IX, haemoglobin alpha locus 1 (HBA1), haemoglobin alpha locus 2 (HBA2), MLH, MSH2, MSH6, apolipoprotein R (APOE), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGT1), amyloid precursor protein (APOE), presentlin-1 (PSENI) and presentilin-2 (PSENI2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolesmia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and
                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDNCA3, melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOR; mismatch repair; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterane; MCPI; syndrome; APP; PSBN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentlin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification -
                                                                                                                                                                                                                                                                                                                                                                                                  Adenosine deaminase deficiency correcting oligo SEQ ID NO: 36.
                                         Length 17;
             1.0%; Score 13.4; DB 1; Length 1: 93.3%; Pred. No. 2.1e+02; mismatches 1; Indels
Sequence 17 BP; 3 A; 3 C; 8 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 7; Page 43; 294pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rice MC;
                                                                                                                                                                                                                                                                         ABA77190 standard; DNA; 17 BP.
                                                                                                                        888 GGAGCTGCGGTACAG 902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-MAR-2000; 2000US-192179P.
01-JUN-2000; 2000US-208538P.
30-OCT-2000; 2000US-244989P.
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                                                                                                                                                       GGAGGTGCGGTACAG 15
                                                                                                                                                                                                                                                                                                                                                        24-JAN-2002 (first entry)
                                                                                 14; Conservative
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                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antilipemic; ss.
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                                                                                                                                                                                                                               RESULT 323
                                                                                 Matches
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The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Pactor VIII, Factor IX, haemoglobin alpha locus 1 (HBA2), MH11, MR22, MK32, MK32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma, APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VI; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein B; LDLR; familial hypercholesterolaemia; UGTI; syndrome; APP; FSENI; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
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various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention.
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                                                                                                                                            Length 17;
                                                                                                                                                                                                     1; Indels
                                                                                                                                            Score 13.4; DB 1;
Pred. No. 2.1e+02;
                                                                                      Sequence 17 BP; 3 A; 8 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                     0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rice MC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABA77193 standard; DNA; 17 BP
                                                                                                                                               1.0%;
                                                                                                                                                                                                                                                              888 GGAGCTGCGGTACAG 902
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27-MAR-2000; 2000US-192179P.
01-JUN-2000; 2000US-208538P.
30-OCT-2000; 2000US-244989P.
                                                                                                                                                                       93.3%;
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                                                                                                                                                                                                                                                                                                           GGAGGTGCGGTACAG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                     14; Conservative
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                                                                                                                                                                             Local Similarity
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                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 324
                                                                                                                                                                                                        Matches
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Gaps

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The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retricoblastoma, BRCA1, BRCA2, CFFR, cyclin-dependent kinase inhibitor 2A (CDKN2A), Apc. Factor V, Factor VII, Factor IX, haemoglobin alpha locus 1 (HBA2), haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6, apolipoprotein E (APOR), LDL receptor (LDIA), upp-glucuronsyltransferase (UGT1), amyloid precursor protein (APC), presentlin-1 (BSBN1) and presentlin-2 (PSBN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent Kinase inhibitor 2A, CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOB; mismatch repair; MSH6; hyperlipidaemia; apolipoprotein B; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSRN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentlin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical
Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonuclectides of the invention.
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                                                                                                                        Length 17;
                                                                                                                                                             1; Indels
                                                                                                                    1.0%; Score 13.4; DB 1;
ilarity 93.3%; Pred. No. 2.1e+02;
Conservative 0; Mismatches 1;
                                                                              Sequence 17 BP; 3 A; 3 C; 8 G; 3 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                             ABA77194 standard; DNA; 17 BP
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01-JUN-2000; 2000US-208538P.
30-OCT-2000; 2000US-244989P.
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                                                                                                                    Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antilipemic; ss.
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                                                                                                                                                                                                                                                                                                      RESULT 325
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The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDKNAA), APC, Factor V, Factor IX, haemoglobin alpha locus apolipoprotein E (APOE), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGT1), amyloid precursor protein (APC), presentlin-1 (PSEN1) and presentlin-2 (PSEN2). These can be used in the gene therapy of diseases
                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; acnoer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MiH1; APOE; mismatch repair; MSH2; MSH6; hypertlipideamia; apolipoprotein E; LULK; familial hypercholesterolaemia; UGT1; syndrome; APP; BSENI; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentin-1; Alzheimer; disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical
haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention.
                                                                                                                                                                   Gaps
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                                                                                                                                Length 17;
                                                                                                                                                                   1; Indels
                                                                                                                               Score 13.4; DB 1;
Pred. No. 2.1e+02;
0; Mismatches 1;
                                                                                           Sequence 17 BP; 3 A; 8 C; 3 G; 3 T; 0 other;
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27-MAR-2000; 2000US-192179P.
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                                                                                                                               1.0%;
Local Similarity 93.3%;
hes 14; Conservative
                                                                                                                                                                                                         888 GGAGCTGCGGTACAG 902
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antilipemic; ss
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01-JUN-2000;
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                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                  RESULT 326
                                                                                                                                                                        Matches
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such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alizheiner's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention.
                                                                                                                                                                   Gaps
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                                                                                                                                   Length 17;
                                                                                                                                                                   1; Indels
                                                                                                                                   Score 13.4; DB 1;
Pred. No. 2.1e+02;
                                                                                                    Seguence 17 BP; 3 A; 2 C; 8 G; 4 T; 0 other;
                                                                                                                                                                    Mismatches
                                                                                                                                                                    0
                                                                                                                                     1.0%;
                                                                                                                         Query Match
Best Local Similarity 93.3%
      88888888
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888 GGAGCTGCGGTACAG 902 2 GGAGGTGCGGTACAG 16 à d

ABA77198 standard; DNA; 17 BP. 24-JAN-2002 (first entry) ABA77198; RESULT 

Adenosine deaminase deficiency correcting oligo SEQ ID NO: 44.

Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein B; LDDR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSEN1; antisense; UDP-glucuronosyltaansferase; amyloid precursor protein; presentlin-1; Albabimer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilipemic; ss

Homo sapiens

WO200173002-A2.

04-OCT-2001

27-MAR-2001; 2001WO-US09761.

27-MAR-2000; 2000US-192176D. 27-MAR-2000; 2000US-192179P. 01-JUN-2000; 2000US-208538P. 30-OCT-2000; 2000US-244989P.

(UYDE ) UNIV DELAWARE

Rice MC Kmiec EB, Gamper HB,

WPI; 2001-639230/73.

Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification -

Claim 7; Page 43; 294pp; English.

The present invention provides single-stranded oligomucleotides which can be used for the targeted alteration of genomic sequences, where the oligomucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDNNAA), APC, Factor V, Factor II, Factor IX, haemoglobin alpha locus (HBA2), MH1, MBH2, MSH2, MSH6, apolloprotein E (APOB), LDL receptor (LDLR), uDP-glucuronosyltransferase (UGI1), amyloid precursor protein (APC), presenilin-1 (PSEN1) and

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presentlin-2 (PSEN2). These can be used in the gene therapy of diseases
                such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention.
          8888888
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Sequence 17 BP; 4 A; 8 C; 2 G; 3 T; 0 other;

Gaps . 0 Length 17; 1; Indels 1.0%; Score 13.4; DB 1; 93.3%; Pred. No. 2.1e+02; ative 0; Mismatches 1; Best Local Similarity 93.3 Matches 14; Conservative Query Match

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GGAGCTGCGGTACAG 902 16 ggaggrgcggracag 2 888 ద à

RESULT 328 ABA80972,

ABA80972 standard; DNA; 17 

BP.

ABA80972;

LDLR mutation correcting oligonucleotide SEQ ID NO: 3818. (first entry) 24-JAN-2002

Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOB; mismarch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein B; LDLR; familial hypercholesterolasemia; UGTI; syndrome; APP; PSEN1; antisense; UDP-glucuromosyltransferase; amyloid precursor protein; presentiin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilipemic; ss

Homo sapiens

WO200173002-A2.

04-OCT-2001.

27-MAR-2001; 2001WO-US09761.

2000US-192176P. 2000US-192179P. 2000US-208538P. 30-OCT-2000; 2000US-244989P. 27-MAR-2000; 27-MAR-2000;

(UYDE ) UNIV DELAWARE.

Gamper HB, Kmiec EB,

WPI; 2001-639230/73.

Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical treating cyst modification

Claim 7; Page 251; 294pp; English.

The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDKN2A) APC, Factor V, Factor VII, Factor IX, haemoglobin alpha locus 1 (HBA1), haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6, apolipoprotein B (APOS), LDL receptor (LDLR), UDP-glucuronosyltransferase

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presenting 1 (PSENZ). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention.
amyloid precursor protein (APC), presenilin-1 (PSEN1) and
     (UGII)
     888888888
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Sequence 17 BP; 6 A; 4 C; 4 G; 3 T; 0 other;

. 0 Gaps .. 0 Score 13.4; DB 1; Length 17; Pred. No. 2.1e+02; 1; Indels 0; Mismatches 745 CATGITGCTGACTIT 759 1.0%; 93.3%; CATGTTGCAGACTTT 1 14; Conservative Query Match Best Local Similarity 15 Matches ð

ABA80973 standard; DNA; 17 BP. ABA80973

ABA80973;

LDLR mutation correcting oligonucleotide SEQ ID NO: 3819.

(first entry)

24-JAN-2002

Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hyperchlesterolaemia; UGTI; syndrome; APP; FSENI; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentilia-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilipemic; ss 

WO200173002-A2.

04-OCT-2001

27-MAR-2001; 2001WO-US09761

27-MAR-2000; 2000US-192176P. 27-MAR-2000; 2000US-192179P. 01-UUN-2000; 2000US-208538P. 30-OCT-2000; 2000US-244989P.

(UYDE ) UNIV DELAWARE

Rice MC

Gamper HB,

Kmiec EB,

WPI; 2001-639230/73.

Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical Claim 7; Page 251; 294pp; English.

The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoplastoma, BRCA1, BRCA2, CFFF, cyclin-dependent kinase inhibitor 2A (CDKWA2A), APC, Factor V, Factor VIII, Factor IX, haemoglobin alpha locus 1 (HBA1), haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6,

Paoni NF; Wood WI;

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UDP-glucuronosyltransferase
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apolipoprotein E (APOE), LDL receptor (LDLR), UDP-glucuronosyltransferas (UGTI), amyloid precursor protein (APC), presentiin-1 (PSENI) and presentiin-2 (PSENI). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophila, hypercholesteroleamia, thalassamia, sitche cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting
                                                                                                                                                                                                                                                                                                                                                                                                                                  cell death, cancer; chromosomal mapping; gene mapping; tissue typing; diagnostic assay; PCR primer; hybridisation; probe; ss.
                                                                                                                                                                              Gaps
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                                                                                                                                                  Query Match
1.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 93.3%; Pred. No. 2.1e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton
Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A,
Grimaldi CJ, Gurney AL, Kljavin IJ, Napier MA, Pan J,
ROY MA, Stewart TA, Tumas D, Watanabe CK, Williams PM,
                                                                                                                                                                                                                                                                                                                                                                                                                     Human; secreted and transmembrane protein; PRO; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                         Human PRO1245 reverse PCR primer SEQ ID NO:493.
                                                                                                                         Seguence 17 BP; 3 A; 4 C; 4 G; 6 T; 0 other;
                                                                                             oligonucleotides of the invention.
                                                                                                                                                                                                                                                                                                          AAF44454 standard; DNA; 17 BP
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; 2000WO-US05004.
; 2000WO-US05841.
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99US-0143048.
99US-0144758.
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2000WO-US00219.
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2000WO-US04341.
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20-MAR-2000; 2000WO-US07377
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                                                                                                                                                                                                                                     3 CATGITGCAGACTIT 17
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24-FEB-2000;
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22-FEB-2000;
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01-DEC-1999;
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us09904568-3.rng

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Generating multiple copies of polynucleotide complementary to RNA, by amplifying RNA with composite primer, second primer, and strand displacement to generate DNA products comprising sequences complementary to RNA.
                                                                                                                                                                                                                                                                                                                                 Nucleic acid sequencing; gene expression; nucleic acid amplification; LGALS1; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                    GALS1 cDNA quantifying primer LGALS13.
                                                                                                                                                                                                                                                                                                                                                                                                                              (NUGE-) NUGEN TECHNOLOGIES INC
                                                                                                                                                                                                                                                                           ABV73116 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                  11-MAR-2002; 2002WO-US07306
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                                                                                                                                                                                                                                                                                                                                                        Synthetic.
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                                                                                                                                                                                             Query Match
Zhang Z;
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                                                                                                                                                                                                            Matches
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(first entry)

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complementary to RNA sequence. The method involves extending first primer hybridized to target RNA (I) with enzyme to form complex having first primer extension product (EPI) and (I), cleaving (I) from the complex, extending second primer hybridized to EPI with enzyme to form complex. Artificial EPI and EP2, and cleaving RNA from the composite primer in the complex. The method is useful for sequencing RNA sequence of interest; for for detecting a mutation in a target RNA using single stranded of a conformation polymorphism; for determining presence or absence of a sequence of interest; for producing a nucleic acid immobilized to a substrate; for characterizing an RNA sequence of interest; for producing a nucleic acid immobilized to a conformation profession profession profession profession profession in a sample; and for preparing a library. The method is also useful for differential amplification of one or more sequence of interest, and for making a cDNA library by preparing a subtractive hybridization probe. Sequences ABV3113-120 represent prime rather cDNA or amplification of the four specific expressed genes in either cDNA or amplification products generated using the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 2 A; 3 C; 6 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the invention.
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                                                                                                                                                                                                                                                                                                                                                                               The present invention describes human secreted and transmembrane PRO proteins. The PRO proteins have cytostatic activity. The PRO proteins can be used for targeted delivery of bioactive molecules, such as toxins, radiolabels or antibodies, that cause cell death. PRO nucleotide sequences, and their fragments, can be used as hybridisation probes, in chromosomal and gene mapping, and in the generation of anti-sense RNA and DNA. They may also be used to produce transgenic animals which are used to develop and screen therapeutically useful reagents. The PRO nucleotide and protein sequence can be used for tissue typing and in treating cancer. Anti-PRO antibodies can be used in diagnostic assays. AAF44270 to AAF44470 represent PCR primers and hybridisation probes used in the isolation of human PRO sequences. AAF44087 to AAF44269 and AAB65164 to AAB65300 represent human PRO polynucleotide and protein sequences given in the exemplification of the present invention.
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                                                                                                                                                                        PRO polynucleotides used to produce polypeptides used to target bioactive molecules such as toxins, radiolabels or antibodies, to specific cells, to cause targeted cell death -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 1 A; 4 C; 7 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                      Example 170; Page 544; 935pp; English.
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                                                                                               WPI; 2001-032160/04.
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                                                                                                                                                                                                                                         Human, gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL -
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0
   Length 17;
                           1; Indels
  1.0%; Score 13.4; DB 1;
93.3%; Pred. No. 2.1e+02;
ive 0; Mismatches 1;
                                                                                                                                                                                                                    Human HTPL scanning oligonucleotide SEQ ID 383.
                                                                                                                                            ABV79137 standard; DNA; 17 BP.
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2001WO-US00665.
2001WO-US00667.
2001WO-US00668.
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2001US-0864761.
2001US-0327898.
                                                       275
Query Match
Best Local Similarity 93.3%;
Matches 14; Conservative
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                                                                                                                                                                                             (first entry)
                                                       261 CCTGGGCTGGCTGAT
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                                                                                                                                                                                                                                                                                                                                                                                                                    30-JAN-2001; 2
30-JAN-2001; 2
30-JAN-2001; 2
30-JAN-2001; 2
30-JAN-2001; 2
23-MAY-2001; 2
09-OCT-2001; 2
                                                                                                                                                                                                                                                                                                                                   EP1229046-A2.
                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001;
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Example 2; Page 114; 718pp; English

The invention relates to generating multiple copies of polynucleotide

Example 5; Page 105; 148pp; English

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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The cotate of Patched, and is a potential tumour suppressor. HTPL is contract in regulating male germ cell development, and the HTPL gene was important in regulating male germ cell development, and the HTPL and in the therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of the HTPL. Such disorders associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and fortal muscle or colon function. HTPL proteins and mucleic acids are clinically useful diagnostic markers and obtenial therapeutic agents for male infertility and cancer. The present oligomucleotide was used in an example from the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 17;
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33.3%; Pred. No. 2.1e+02;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human HTPL scanning oligonucleotide SEQ ID 384.
                                                                                                                                                                                                                                                                                                                                                                                                             Seguence 17 BP; 3 A; 6 C; 7 G; 1 T; 0 other;
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2001WO-US00665.
2001WO-US00667.
2001WO-US00668.
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2001US-0864761.
2001US-0327898.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93.3%;
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30-JAN-2001;
30-JAN-2001;
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23-MAY-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV'8759 to ABV'8762 and ABB98519 to ABB98520). HTPL brotein (HTPL, see ABV'8759 to ABV'8762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair diangerences between the two isoforms with a few single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL codon in HTPL-S (S for short) compared to HTPL-S a role similar to that of Patched, and is a potential tumour suppressor. HTPL is comportant in regulating male germ cell development, and the HTPL gene was useful for diagnosing a disorder caused by mutation in HTPL, and in company and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of such disorders associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and HTPL. Such disorders include disorders of testis, or adrenal, adult and skeletal muscle or colon function. HTPL proceins and mucleic acids are clinically useful diagnostic markers and potenial therapeutic acids are colon function and potenial therapeutic acids are male infertility and cancer. The present oligonucleotide was used in an example from the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  / Match 1.0%; Score 13.4; DB 1; Length 17; Local Similarity 93.3%; Pred. No. 2.1e+02; nes 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human HTPL scanning oligonucleotide SEQ ID 385.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 4 A; 5 C; 7 G; 1 T; 0 other;
Example 2; Page 114; 718pp; English.
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2001US-0864761.
2001US-0327898.
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2001WO-US00665.
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30-JAN-2001; 2
23-MAY-2001; 2
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30-JAN-2001;
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Matches
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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL brack will be a single base pair differences between the two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-8 (8 for short) compared to HTPL-10 (L for long). HTPL shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The content of Patched, and is a potential tumour suppressor. HTPL is similar to that of Patched, and is a potential tumour suppressor. HTPL is see a mapped to human chromosome 10pl2.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of therapy and cancer include disorders of testis, or adrenal, adult and the HTPL. Such disponsition arkers and potential therapeutic acids are colinically useful diagnostic markers and potential therapeutic acids are male infertility and cancer. The present oligonucleotide was used in an example from the invention.
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Best Local Similarity 93.3%; Pred. No. 2.1e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
partners, and for treating subjects having defects in HTPL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 17 BP; 5 A; 4 C; 7 G; 1 T; 0 other;
                                        Example 2; Page 114; 718pp; English.
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2001WO-US00669.
2001WO-US00670.
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2001WO-US00665.
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10-OCT-2001; 2001US-0328205
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABB83999), a sequence having 65% sequence of 6711, colling to (SI), and the sequence comprising at least 8 contiguous amino or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interactes with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) conceded by altered expression of human POSHL1 including disease and treating caused by altered expression of human POSHL1 including diseases and (II) is treating cancer, they useful in the development of vaccines and (II) is useful for measuring and for suverying gene expression and creating are useful for measuring and for suverying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; POSH1 1; SH3 domain; POSH-like signalling protein 1; oncogene;
Rho GTPase; signal transduction; gene expression; cancer; vaccine;
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       Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
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                                                                                  Example 2; SEQ ID NO 494; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human POSHL1 scanning oligonucleotide SEQ ID NO 495.
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30-JAN-2001; 2001WO-US00666.
30-JAN-2001; 2001WO-US00667.
30-JAN-2001; 2001WO-US00669.
30-JAN-2001; 2001WO-US00669.
20-JAN-2001; 2001WO-US00669.
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ABV89782 standard; DNA; 17 BP.
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nes 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                     the invention.
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30-JAN-2001; 2001WO-US00666.
30-JAN-2001; 2001WO-US00667.
30-JAN-2001; 2001WO-US00668.
30-JAN-2001; 2001WO-US00669.
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                            Shannon M;
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RANGE STATE 
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The invention relates to an isolated SH3 domain (POSH)-like signalling correctin 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABB83999), a sequence having 65% sequence identity to (SI), (SI) having 95% deviations, a sequence comprising a sequence identity to (SI), (SI) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small Grases as well as downstream components of the signal transduction pathway. (I) is useful as concoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and creating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which care useful for measuring and for surveying gene expression and creating cransgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the measuring cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of the invention.
Note: The present sequence did not form part of the printed
specification, but is based on sequence information supplied to Derwent
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                                                                                                                                                                                                                                Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Match 1.0%; Score 13.4; DB 1; Length 17; Local Similarity 93.3%; Pred. No. 2.1e+02; es 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                         Example 2; SEQ ID NO 496; 60pp + Sequence Listing; English.
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23-MAY-2001; 2001US-0864761.
10-OCT-2001; 2001US-0328205.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15 crcccedaccrcrec 1
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                                                                                                                                                                                   WPI; 2002-684061/74.
                                                                          (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABV90552;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         à
                                                                                                                                                                                                                                                                            The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABBB3999), a sequence having 65% sequence of 730 amino (CI) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acida. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small Grases as well as communitying a specific binding partner. (I) and mucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and oreating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to Derwent by the European Patent Office.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, POSHL 1; SH3 domain, POSH-like signalling protein 1, oncogene, Rho GTPase; signal transduction, gene expression, cancer, vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                               Novel human SH3 domain. (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.1e+02; rative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                  Example 2; SEQ ID NO 495; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human POSHL1 scanning oligonucleotide SEQ ID NO 496.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 4 A; 4 C; 8 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABV89783 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene therapy; transgenic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     631 CTCCAGGAGCTCTGC 645
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CICCCGGAGCICIGC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-DEC-2002 (first entry)
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Gaps

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30-JAN-2001; 2001MO-US00663. 30-JAN-2001; 2001MO-US00664. 30-JAN-2001; 2001MO-US00665. 30-JAN-2001; 2001MO-US00666.

28-JAN-2002; 2002EP-0001165.

11-SEP-2002.

99US-123972P. 99US-133459P. 99US-140650P. 99US-144653P. 99US-144758P.

99US-146222P. 99US-149395P. 99US-151689P. 99WO-US20111.

99WO-US21090. 99WO-US28313. 99WO-US28634 2000WO-US00219

us09904568-3.rng

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The present invention relates to the isolation of novel human PRO polypeptides (AAU86128-AAU86162) and the polymucleotide sequences encoding them. The PRO polypeptides, agonists, antagonists or anti-PRO antibodies are useful for treating benign or malignant tumours (e.g. renal, kidney, bladder, breast, etc), leukaemias and lymphoid malignancies, other disorders such as neuronal, glial, astrocytal, hypothalamic, glandular, macrophagal, stromal and blastocoelic disorders, inflammatory, immune and angiogenic disorders. The polymucleotide sequences are also useful in gene therapy. The present sequence represents a PCR primer used in the methods of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Thirty five nucleic acids encoding PRO polypeptides, useful for treating benign or malignant tumours, leukaemias and lymphoid malignancies, inflammatory, angiogenic and immunologic disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 1 A; 4 C; 7 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 26; Page 143; 302pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-205567/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ashkenazi AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Marsters SA,
Watanabe CK,
                                                                                                                                                                                                                                                                                                                                                                                                                 05-JAN-2000;
                                                                                                                                       22-JUN-1999;
                                                                                                                                                                                                               28-JUL-1999;
                                                                                                                                                                                                                                        17-AUG-1999;
                                                                                                                                                                                                                                                               31-AUG-1999;
                                                                                                                                                                                                                                                                                                                15-SEP-1999;
                                                                     11-MAY-1999;
                                                                                       02-JUN-1999
                                                                                                                                                                                        26-JUL-1999
                                                                                                                                                                                                                                                                                                                                            30-NOV-1999,
                                                                                                                                                                                                                                                                                                                                                                 01-DEC-1999
The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino acids (21, ABB83999), a sequence having 54 sequence of 631), (S1) having 954 deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GrPases as well as downstream components of the signal transduction pathway. (I) is useful of for identifying a specific binding partner. (I) and nucleic acids (II) concounting (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and treating caused by altered expression of human POSHL1 including diagnosing and treating caused by measuring and for surveying gene expression and creating are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to Derwent by the European Patent Office.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
                                                                                                                                                                                                                                                                                                           Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.1e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; SEQ ID NO 1265; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 4 A; 7 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    263 TGGGCTGGCTGATCA 277
                    30-JAN-2001; 2001WO-US00668.
30-JAN-2001; 2001WO-US00669.
30-JAN-2001; 2001WO-US006670.
323-MAY-2001; 2001US-0864761.
  2001WO-US00667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 93.3
Matches 14; Conservative
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                                                                                                                                                                                                                                                                       WPI; 2002-684061/74.
                                                                                                                                                                        (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of the invention.
                                                                                                                                                                                                                         Shannon M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16
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Hillan KJ;

Gurney AL, Smith V,

Goddard A, Godowski PJ, Pan J, Pitti RM, Roy MA,

Pan J, E Wood WI;

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Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic; antiinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinetic;
                               Gaps
                               ;
0
1.0%; Score 13.4; DB 1; Length 17; 13.3%; Pred. No. 2.1e+02;
                               1; Indels
                                                                                                                                                                                                                                                          Human CLCAl gene enzymatic nucleic acid #1062.
                               0; Mismatches
                                                                                                                                                                      ABK56691 standard; RNA; 17 BP
 Ouery Match 1.0%;
Best Local Similarity 93.3%;
                                                               494 GIGIGCAGCGICTIG 508
                                                                                           erececeacina 15
                                                                                                                                                                                                                                (first entry)
                                 14; Conservative
                                                                                                                                                                                                                                                                                                                                                         acetylcysteine.
                                                                                                                                                                                                                                  02-JUL-2002
                                                                                                                                                                                                    ABK56691;
                                                                                                                                         RESULT 340
                                  Matchea
                                                                                                                                                           ABK56691
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Reverse PCR primer #2 for gene amplification analysis of human PRO1245.

(first entry)

15-JUL-2002

ABK40398;

ABK40398 standard; DNA; 17 BP

RESULT 339 ABK40398

ð g Human; PRO; benign tumour; malignant tumour; lymphoid malignancy; leukaemia; neuronal disorder; stromal disorder; blastocoelic disorder; inflammatory disorder; immune disorder; angiogenic disorder; cytostatic; neuroprotective; PCR; primer; ss.

Homo sapiens

11-FEB-2000; 2000WO-US03565

WO200153486-A1. Homo sapiens.

26-JUL-2001

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Enzymatic polynuclectide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma
                                                                                          McSwiggen J, McKenzie T, Ayers D, Szymkowski DE;
                                                                                                                                                           Claim 4; Page 78; 152pp; English.
                              09-AUG-2001; 2001WO-US24970.
                                            09-AUG-2000; 2000US-224383P
                                                           (RIBO-) RIBOZYME PHARM INC.
                                                                                                               WPI; 2002-217145/27.
                                                                          (THOM/) THOMPSON J.
WO200211674-A2.
               14-FEB-2002.
                                                                                         Thompson J,
                                                                                                 Grupe A;
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The invention relates to enzymatic nucleic acid molecules that down regulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic fibrosis, obstructive pulmonary disease (CCDPD), chronic bronchitis, asthma, cystic that are related to or will respond to the levels of CLCA1 in a cell or hence, are useful for treatment of a patient having a condition associated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the present the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapy, bronchodilators, corticosteroids, nucleic acids of the invention are also used as diagnostic agents. The examine genetic drift and mutations within diseased calls or to detect the presence of CLCAI RNA in a cell. This sequence represents an enzymatic nucleic acid molecule of the invention. Gaps ö 1.0%; Score 13.4; DB 1; Length 17; 80.0%; Pred. No. 2.1e+02; ative 2; Mismatches 1; Indels Sequence 17 BP; 7 A; 6 C; 2 G; 2 U; 0 other; 646 ATCCCCCAAGACCTG 660 Query Match
Best Local Similarity 80.05
Matches 12; Conservative

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2 Auccaccaadaccud 16

ABK18409 standard; RNA; 17 BP. (first entry) 09-APR-2002 ABK18409; RESULT 341 ABK18409 

Human ERG hammerhead ribozyme target sequence, Seq ID No 1056.

Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular Jaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;

Homo sapiens

WO200188124-A2.

22-NOV-2001.

16-MAY-2001; 2001WO-US15866.

16-MAY-2000; 2000US-0572021.

(RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.

Randi AM: Von Carlowitz I, McSwiggen JA, Mclaughlin F, WPI; 2002-082995/11. Jarvis T,

Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome

Claim 4; Page 78; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (EEG). (I) is useful for treating conditions selected from cancer. lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, tumour angiogenesis, diabetic retinopathy, macular degeneration, vertuca tumour angiogenesis, diabetic retinopathy, macular degeneration, vertuca cumour glaucoma, Wippel Trenaunay-Weber Syndrome, Osie-Weber-rendu CC wedgaris, angiofibroma of tubercous sclerosis, port-wine stains, Sturge syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of EEG, by contacting cells of the patient with (I) under conditions suitable for under conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a coll.) by contacting the cell with (I). (I) is useful for clasving sample coll to diseases related to the expression of ERG, and as diagnostic tool tool context of the presence of ERG RNA in a cell. (I) is useful for specifically catamine genes that share homology with ERG gene or ERG fusion genes.

Cation such as MSZ2719 represent nucleic acids, including antisense and cargred enzymatic nucleic acid molecules which regulate expression of ERG, and enzymatic nucleic acid molecules which regulate expression of ERG, and cargrant enzymatic nucleic acid molecules which regulate expression of ERG, and enzymatic nucleic acid molecules.

Sequence 17 BP; 6 A; 5 C; 2 G; 4 U; 0 other; 

.; 0 1.0%; Score 13.4; DB 1; Length 17; 30.0%; Pred. No. 2.1e+02; ive 2; Mismatches 1; Indels 80.08; Best Local Similarity 80.0 Matches 12; Conservative Query Match

.; 0

Gaps

UNAUACCAAACCCAG 17 21 TTAAACCAAACCCAG 35 m

à 엄

ABK18410 standard; RNA; 17 09-APR-2002 (first entry) ABK18410; ABK18410 

RESULT 342

ВЪ.

Human ERG hammerhead ribozyme target sequence, Seq ID No 1057.

Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis;

neovascular glaucoma, myopic degeneration, arthritis, verruca vulgaris; angiófibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Leukaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme; tumour angiogenesis; diabetic retinopathy; macular degeneration;

Homo sapiens.

WO200188124-A2

22-NOV-2001

16-MAY-2001; 2001WO-US15866.

16-MAY-2000; 2000US-0572021

(RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.

Jarvis T, Von Carlowitz I,

WPI; 2002-082995/11.

Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome

Claim 4; Page 78; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating carbons and conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, or ungarial, angiofibroma of tuberous sclerosis, port-whose stains, Sturge Neber syndrome, Rippel-Trenaunay-Weber syndrome, Osler-Weber-rendu Syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. Euckeamia or tumour angiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or coll, by contacting the cell with (I) (I) is useful for cleaving RNA of ERG gene, by contacting the cell with (I) (I) is useful for cleaving RNA of ERG gene, by contacting the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect conference of ERG RNA in a cell. (I) is useful for specifically transpared mutations within diseased cells or to detect angesting genes that share homology with ERG gene or ERG fusion genes. ABK/1354-ABK/2719 represent nucleis added molecules which regulate expression of ERG, and exargmatic nucleic acide molecules which regulate expression of ERG, and related PCR primers of the invention. 

Sequence 17 BP; 6 A; 5 C; 2 G; 4 U; 0 other;

ö Length 17; 1; Indels Query Match 1.0%; Score 13.4; DB 1; Best Local Similarity 80.0%; Pred. No. 2.1e+02; Matches 12; Conservative 2; Mismatches 1;

0

Gaps

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RESULT 343

ABK19335,

ABK19335 standard; RNA; 17 BP. ABK19335 EXE EXE

09-APR-2002 (first entry)

Human ERG Amberzyme target sequence Seq ID No 1982.

Human, hammerhead ribozyme, cytostatic, antitumour; antidiabetic, ophthalmological; antiatthritic; antipscriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; pscriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; necowascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme; amberzyme.

Homo sapiens.

WO200188124-A2. 

22-NOV-2001.

16-MAY-2001; 2001WO-US15866.

Randi AM;

Mclaughlin F,

McSwiggen JA,

16-MAY-2000; 2000US-0572021.

(RIBO-)

RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LID. Von Carlowitz I, McSwiggen JA, Mclaughlin F, Randi AM; WPI; 2002-082995/11. Jarvis T,

Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome

Claim 4; Page 126; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, metlanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, tumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, nervascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca vulgaris, angiogenesis, stopel-Trenaunay-Weber syndrome, Osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies the treatment. Includacing suitable for the treatment. Includacing or under conditions suitable for the treatment. Leukacmia or tumour angiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting the cell with (I). (I) is useful for cleaving RNA of ERG gene, by contacting (I) with RNA, in the presence of a divalent confinesses related to the expression of ERG such or to detect the presence of ERG RNA in a cell. (I) is useful for specifically confinesses that share homology with ERG gene or ERG fusion genes. ABMI1354-ABK22719 represent nucleic acide, including antisense and cerametric mucleic acid molecules which regulate expression of ERG, and cerametric mucleic acid molecules which regulate expression of ERG, and cerametric mucleic acid molecules which regulate expression of ERG, and cerametric mucleic acid molecules which regulate expression of ERG, and cerametric molecules which regulate expression of ERG, and cerametric molecules and cerametric mucleic acid molecules acid including antisense and cell molecules which regulate expression of ERG, and and cell molecu

Sequence 17 BP; 7 A; 1 C; 6 G; 3 U; 0 other;

Gaps .. 0 1.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.1e+02; tive 0; Mismatches 1; Indels 14; Conservative Query Match Best Local Similarity Matches

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TTAGCATACTCCTCA 62 48

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16 TTAGCATCCTCA 2
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RESULT 344

ABT39230.standard; DNA; 17 BP. ABT39230;

(first entry) 12-JUN-2003 

Tumour suppression related human fukutin oligo SEQ ID No 4867.

Cytostatic, virucide, neuroprotective, nootropic, neuroleptic, gene chip, antisense, sense, tumour, cell degeneration, cancer, Alzheimer's disease; schizophrenia, protein chip, gene therapy, tumour suppression, human fukutin, ds.

Homo sapiens

WO2003025175-A2.

27-MAR-2003.

17-SEP-2002; 2002WO-IB04208.

17-SEP-2001; 2001FR-0011978.

(MOLE-) MOLECULAR ENGINES LAB.

Felerman A, Amson R,

Tuijnder M;

WPI; 2003-313353/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells

Disclosure; Page 603; 720pp; French

The invention relates to a novel isolated 17 mer mucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, polypeptides, vectors containing the nucleic acids, polypeptides, vector or antibodies directed against the polypeptides are useful for proparation of pharmaceuticials for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and any any of the life and any or any any of the supression of the 17 mer nucleic acids in a striant paramles is nearly the expression of the 17 mer nucleic acids in a striant paramles is nearly the expression of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polymucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention.

Sequence 17 BP; 8 A; 3 C; 4 G; 2 T; 0 other;

. 0 1.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.1e+02; ive 0; Mismatches 1; Indels Local Similarity ses 14; Conserv Query Match Best Loca Matches

à Db

ABT39985 standard; DNA; 17 BP. 

ABT39985;

(first entry) 13-JUN-2003 Tumour suppression related human fukutin oligo SEQ ID No 5622.

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antieense; sense; tumour; cell degeneration; cancer; Alzhaimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.

Homo sapiens

WO2003025175-A2.

27-MAR-2003.

17-SEP-2002; 2002WO-IB04208.

17-SEP-2001; 2001FR-0011978.

(MOLE-) MOLECULAR ENGINES LAB

Tuijnder M; relerman A, Amson R,

WPI; 2003-313353/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells

Disclosure; Page 691; 720pp; French.

The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding NNW. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the peparation of plarmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell cleaneration, specifically cancer but also Albreimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for these patients and the containing the containing the patient samples is useful for diagnosis and/or progness and seconds. diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention.

Sequence 17 BP; 2 A; 10 C; 3 G; 2 T; 0 other;

0 1.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.1e+02; ive 0; Mismatches 1; Indels 14; Conservative Best Local Similarity Matches 14; Conserv Query Match

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Gaps

760 CGGTGGCGGGTGGAT 774 16 cécacécederécar

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Gaps

RESULT 346

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05-JUN-1998;
09-JUN-1998;
10-JUN-1998;
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hypertrophy of neonatal heart; angiogenesis; wound healing;
                                                                          cardiac insufficiency disorder; cancer; tumour; immune response; adrenal cortical capillary endochelial growth; c-fos induction; vascular endothelial growth factor inhibition; VEGF inhibition; endothelial cell growth inhibitor; T-lymphocytes stimulation; retinal neurons cell survival; rod photoreceptor cell survival; retinal disorder; retinitis pigmentosum, kidney disorder; mammalian kidney mesangial cell proliferation; Berger disease; dermatitis; herpetiformis; Crohn's disease; chondrocyte proliferation; chondrocyte redifferentiation; sports injury; arthritis; gene; ds.
                                                     Novel human secreted or transmembrane protein PRO183 DNA
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2001WO-US21066.
2001WO-US21735.
97US-049787P.
97US-062250P.
        ABX80463 standard; DNA; 17 BP
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2000WO-US07377.
2000WO-US08439.
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98WO-US19330.
98WO-US19437.
98WO-US21141.
99WO-US25108.
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2000WO-US04914.
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2000WO-US13705.
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2000WO-US14941.
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2000WO-US23328
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2000WO-US20710
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99WO-US21090
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                                     (first entry)
                                                                                                                                                                       US2002132252-A1,
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02-MAR-2000;
15-MAR-2000;
20-MAR-2000;
30-MAR-2000;
15-MAY-2000;
17-MAY-2000;
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11-FEB-2000;
18-FEB-2000;
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20-DEC-1999;
06-JAN-2000;
                                                                                                                                                         Homo sapiens.
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24-FEB-2000;
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28-JUL-2000
                                                                    Human; PRO;
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                                     28-APR-2003
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                      ABX80463;
 ABX80463
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The invention describes an isolated human PRO polypeptide. The PRO polypeptides are useful in detecting PRO polypeptides in a sample, in linking a bioactive molecule to a cell expressing a PRO polypeptide, and in modulating at least one biological activity of a cell expressing a PRO polypeptide. PRO1312 stimulates hypertrophy of neonatal heart and is thus useful for treating cardiac insufficiency disorders. PRO1154 and PRO1186
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Wood WI;
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Perzara N, Rong S, Gerber H, Gerritsen ME, Goddard A,
Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J,
Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM,
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98US-088167P.
98US-088212P.
98US-088212P.
98US-088217P.
98US-088517P.
98US-088734P.
98US-088742P.
9705-065311P.
9708-065710P.
9808-075945P.
9808-083322P.
9808-08760P.
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9808-086026P.
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98US-088858P.
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98US-089440P.
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98US-089908P
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stimulate adremal cortical capillary endothelial growth, and PRO536, PRO943, PRO828, PRO168 or PRO535, PRO943, PRO828, PRO168 or PRO535, PRO943, PRO828, PRO168 or PRO535, PRO943, PRO943, PRO1126, PRO160 and PRO187, induce c-fos in endothelial cells, and are thus cueful for treating conditions or disorders where angiogenesis would be beneficial, e.g. wound healing and antagonist of this polypeptide are useful for treating conditions two stimulated this polypeptide are endothelial growth factor (VEGF) Stimulated proliferation of endothelial growth factor inhibiting endothelial cell growth in mammals which would be beneficial in inhibiting tumour growth. PRO826, PRO1068, PRO1184, PRO1346 and PRO1135 stimulate proliferation of stimulated T-lymphocytes and are therapeutically useful for enhancing immune response. PRO826, PRO108 or PRO1132 enhance survival/proliferation of retinal neurons cells (PRO1132 is also enhances survival/proliferation of disorders of injuries, e.g. retinitis pigmentosum, AMD. PRO819, PRO813 and therefore are useful for treating kidney disorders associated with dermaltian kidney mesangial cells, and therefore are useful for treating kidney disorders associated with dermaltis, herpetiformis or Crohn's disease. PRO110, PRO844, PRO112, PRO1192 and PRO1187 induce the comprehence are thus useful for treating sports and PRO1186. This care thus useful for treating sports and arthritis. This care thus useful for treating sports in polymucleotide.
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                                                                                                                                                                                                                                                                                                                                 Query Match 1.0%; Score 13.4; DB 1; Length 17; Best Local Similarity 93.3%; Pred. No. 2.1e+02; Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human secreted/transmembrane protein, TaqMan probe #22.
                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 1 A; 4 C; 7 G; 5 T; 0 other;
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98WO-US25108.
99WO-US00106.
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98WO-US19437.
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01-DEC-1999;
01-DEC-1999;
16-DEC-1999;
20-DEC-1999;
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02-JUN-1999;
15-SEP-1999;
15-SEP-1999;
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01-DEC-1998;
05-JAN-1999;
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17-SEP-1998;
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96US-088030P.
98US-088033P.
98US-088157E.
98US-08812P.
98US-088512P.
98US-08855P.
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22-FEB-2000;
24-FEB-2000;
24-FEB-2000;
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28-FEB-2001;
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04-JUN-1998;
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Human; PRO; hypertrophy of neonatal heart; angiogenesis; wound healing; cardiac insufficiency disorder; cancer; tumour; immune response; adrenal cortical capillary endcthelial growth. c-fos induction; vascular endchelial growth factor inhibition; VEGF inhibition; endchelial cell growth inhibitor; T-lymphocytes stimulation; retinal heurons cell survival; rod photoreceptor cell survival; retinal disorder; retinits pjementosum; kidney disorder; mammalian kidney mesangial cell proliferation; Berger disease; dermatitis; herpetiformis; Crohm's disease; chondrocyte proliferation; chondrocyte redifferentiation; sports injury; arthritis; gene; ds.
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         98US-097218P.
98US-097661P.
98US-097952P.
98US-097954P.
98US-097971P.
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98WO-US19330.
98WO-US19437.
98WO-US25108.
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98US-098014P.
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20-DEC-1999; 99WO-US30911.
05-JAN-2000; 2000WO-US00219.
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17-SEP-1998;
07-OCT-1998;
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05-JAN-1999;
08-MAR-1999;
         20-AUG-1998
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ABX81350
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PR 10-PED-2000, 2000WO-JUS0541.

PR 10-PED-2000, 2000WO-JUS054.

PR 10-PED-2000, 2000WO-JUS054.
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Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Fong S, Gebber H, Gerritsen ME, Goddard A, Godowski PJ; Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J, Paoni NF; Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI; Zhang Z;
02-JUN-2000; 2000WO-USI5264.
28-JUL-2000; 2000WO-US20710.
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Pred. No. 2.1e+02;
0; Mismatches 1; Indels
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2000WO-US03565.
2000WO-US04341.
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Matches 14; Conservative
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ABX90440
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22-MAY-2000;
30-MAY-2000;
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  The invention discloses isolated PRO secreted/transmembrane polypeptides comprising a sequence without signal peptide and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. The PRO polypeptides or polympetides are also useful in gene therapy, in chromosome dentification, as chromosome markers, or in generating probes. The PRO polympetides are useful as molecular markers for protein properties are useful as molecular markers. The PRO polypeptides and nucleic acids may also be used in tissue typing, Anti-PRO antibodies are useful in diagnostic assays for PRO, and in affinity purification of PRO from recombinant cell culture or natural sources. The sequences presented in ABX90083-ABX90468 are the genes encoding, the primers amplifying and the probes detecting the PRO polymucleotides of the invention.

Probes detecting the PRO polymucleotides of the invention.

Commat from USPTO at seqdata.uspto.gov/sequence.html.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; PRO; PCR; ss; cytostatic; tumour; cancer; breast; lung; stomach; liver; horse; cow; dog; cat; sheep; pig; goat; rabbit; ADEPT; primer; antibody-dependent enzyme mediated prodrug therapy.
New transmembrane polypeptides and nucleic acids encoding the polypeptides, useful in gene therapy, in chromosome identification, as chromosome markers, or in generating probes -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 1 A; 4 C; 7 G; 5 T; 0 other;
                                                                                          Example 170; Page 297; 650pp; English.
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98WO-US25108.
99WO-US00106.
99WO-US05028.
99WO-US12252.
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99WO-US30095.
99WO-US30911.
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98WO-US19437
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99WO-US21547
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99WO-US28301
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02-JUN-1999;
15-SEP-1999;
15-SEP-1999;
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01-DEC-1999;
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16-DEC-1999;
20-DEC-1999;
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Human; probe; ss; PRO; secreted; transmembrane; signal peptide; pharmaceutical; diagnostic; biosensor; bioreactor; tumour; therapeutic; colon cancer; lung cancer; breast cancer; cancer; gene therapy; TaqMan.
                                                                                                                                                                                                                                                                                                                                                                               Human sectreted/transmembrane protein, TaqMan probe #22.
                                                                                                                                                                                                      1.0%; Score 13.4; DB 1;
Local Similarity 93.3%; Pred. No. 2.1e+02;
les 14; Conservative 0; Mismatches 1;
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99WO-US0508
99WO-US05028
99WO-US21252
99WO-US21547.
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2000WO-US00219.
2000WO-US00376.
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2000WO-US04341.
2000WO-US04414.
2000WO-US04914.
98US-097141P.
98US-09761P.
98US-097652P.
98US-097954P.
98US-097954P.
98US-097974P.
98US-097974P.
98US-097978P.
98US-097978P.
98US-097978P.
98US-097978P.
98US-097978P.
98US-10986P.
98US-100858P.
98US-112396P.
99US-112396P.
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2000WO-US05841
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15-SEP-1999;
30-00V-1999;
01-DEC-1999;
01-DEC-1999;
05-JAN-2000;
06-JAN-2000;
11-FEB-2000;
22-FEB-2000;
24-FEB-2000;
19-AUG-1998;
24-AUG-1998;
24-AUG-1998;
26-AUG-1998;
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11-SEP-1998;
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17-SEP-1998;
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02-MAR-2000;
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Best Local S:
Matches 14
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ABX79647
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   98US-089653E-
98US-089901P-
98US-089901P-
98US-089908P-
98US-089948P-
98US-090254P-
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98US-090444P-
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98US-0904435P-
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98US-091818P-
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980S-096773P
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980S-09697P
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17-AUG-1998;
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04-AUG-1998
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Gaps

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Length 17; 1; Indels Eaton DL;

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98US-088824P.
98US-088826P.
98US-08861P.
98US-08861P.
98US-089105P.
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98US-08759P.
98US-088021P.
98US-088025P.
98US-088026P.
98US-088028P.
98US-088038P.
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98US-089598P.
98US-089599P.
98US-089600P.
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970S-065311P.
970S-066770P.
980S-075945P.
980S-078910P.
980S-083322P.
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2001WO-US21735.
97US-049787P.
97US-062250P.
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98US-087106P.
98US-087607P.
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2000WO-US20710.
2000WO-US22031.
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2000WO-US23328.
2000WO-US30952.
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2001WO-US06520.
2001WO-US17800.
     2000WO-US06884.
2000WO-US07377.
2000WO-US08439.
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29-JUN-2001;
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17-MAY-2000;
22-MAY-2000;
30-MAY-2000;
02-JUN-2000;
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01-DEC-2000;
28-FEB-2001;
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16-JUN-1998;
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20-MAR-1998;
28-APR-1998;
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28-MAY-1998;
02-JUN-1998;
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04-JUN-1998;
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24-AUG-2000;
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04-JUN-1998;
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The invention discloses isolated PRO secreted/transmembrane polypeptides comprising a sequence without signal peptide and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that encoding them. The polypeptide, for linking a bicactive encoding them. The properties, for linking a bicactive molecule to a cell expressing a PRO polypeptide, for linking a bicactive molecule to a cell expressing a PRO protein and for modulating at least on the properties of the properties of the properties of the properties of conclusivity of a cell. The PRO polypeptides or polynucleotides are also useful as pharmaceuticals, diagnostics, biosensors or bioreactors, for detecting or treating e.g. tumours in mannals, e.g. humans, dogs, cats, cattle, horses, sheep, pigs, goats or rabbits as targets for therapeutic intervention in certain cancers (e.g. colon, lung or breast cancers) and diagnostic determination of the presence of these cancers. The PRO polypeptides are also useful as molecular weight markers or for for homosome identification. The PRO genes are useful as hybridisation probes or for screening libraries of human cond. Agnowic or mRNA, The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequences presented in ABX79290-ABX79675 are the genes encoding, the primers among the primers of the propes detecting the PRO polynucleotides of the
           Godowski PJ;
Paoni NF;
Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Note: The sequence data for this patent is also available in electronic format from USPTO at segdata.uspto.gov/sequence.html.
                                                                                                                                                                       New secreted and transmembrane PRO polypeptides (e.g. PRO183, PRO184, PRO361 or PRO846) useful as targets for therapeutic intervention in cancers (e.g. lung or breast cancers), or for diagnosing these cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton Persrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J, Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seguence 17 BP; 1 A; 4 C; 7 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                             Example 170; Page 294; 647pp; English.
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06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention.
                                                                 Roy MA,
Zhang Z;
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ABZ60185/c
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acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2. K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HHV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ56989 - ABZ65151, ABZ64544 - ABZ65531, ABZ66520 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention.
                                                                                                                                                                                                                        The invention relates to a novel short interfering RNA (siRNA) nucleic
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                                                                                                                 Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.1e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 4 A; 5 C; 2 G; 6 U; 0 other;
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                                                                                                                                                                                           Claim 58; Page 90; 185pp; English.
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06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
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                (RIBO-) RIBOZYME PHARM INC.
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Best Local Similarity 93.3
Matches 14; Conservative
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                                                   Mcswiggen J;
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                                     invention relates to a novel short interfering RNA (siRNA) nucleic
                                                    acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HRR2, K-RRS, H-RAS, N-Rhuman immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HRR2, K-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in RBZ59899 - ABZ65216, ABZ65231, ABZ6520 - ABZ65216, ABZ65531, sequences for the human ribozymes of the invention.
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enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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1.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 2.1e+02;
Matches 14; Conservative 0; Mismatches 1;
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Claim 58; Page 109; 185pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                        281 AGGAAGCAGCAAA 295
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06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
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                                                                                                                                                                                                                                                                                                                                          Query Match
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acid molecule or an enzymatic nucleic acid molecule, that modulates burnession of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule invention has cyrostatic, anti-HHV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, The sequences shown in ABZS989 - ABZ66216, ABZ66531, ABZ66530 - ABZ66531, ABZ66531 - ABZ66531, ABZ66530 - ABZ66531, ABZ66531 - ABZ66531 in the invention.
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The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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                                                                                                             Query Match 1.0%; Score 13.4; DB 1; Length 17; Best Local Similarity 93.3%; Pred. No. 2.1e+02; Matches 14; Conservative 0; Mismatches 1; Indels
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                                                                            Sequence 17 BP; 2 A; 4 C; 8 G; 3 U; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Human H-Ras DNAzyme target #694.
                                                                                                                                                                                       1293 TGCTCAGCCTGGCCC 1307
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06-JUN-2001; 2001US-296249P.
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Human; PRO polypeptide; secreted protein; transmembrane protein;
genetic disorder; antibacterial; immunosuppressive; PCR; primer; ss.
                                                                                               Human PRO DNA PCR primer #125
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97US-065311P.
97US-066770P.
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99WO-US00106.
99WO-US05028.
99WO-US12252.
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24-AUG-2000; 2000WO-US23328
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2001US-0989731
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                                                ABX64286 standard; DNA; 17
                                                                               26-FEB-2003 (first entry)
                                                                                                                                                      US2002103125-A1.
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11-FEB-2000;
18-FEB-2000;
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24-FEB-2000;
24-FEB-2000;
02-MAR-2000;
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28-FEB-2001;
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                                                                                                                                      Homo sapiens.
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20-MAR-2000;
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15-SEP-1999;
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30-NOV-1999;
01-DEC-1999;
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05-JAN-1999
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                                                                ABX64286;
                                RESULT 356
                                         ABX64286
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Query Match 1.0%; Score 13.4; DB 1; Length 17; Best Local Similarity 73.3%; Pred. No. 2.1e+02; Matches 11; Conservative 3; Mismatches 1; Indels

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Gaps

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Godowski PJ;
Paoni NF;
Wood WI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel secreted and transmembrane polypeptide for modulating biological activity of cell expressing the polypeptide, identifying agonists or antagonists of polypeptide, and as molecular weight markers -
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Berrara M, Rong S, Gerber H, Gerritsen ME, Goddard A,
Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J,
Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM,
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98US-075945P.
98US-078910P.
98US-083321P.
98US-08450P.
98US-08160P.
98US-087607P.
98US-087607P.
98US-087607P.
98US-088025P.
98US-088025P.
98US-088023P.
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98US-088023P.
98US-08812P.
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98US-088174P.
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98US-089801P.
98US-089907P.
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04-JUN-1998;
04-JUN-1998;
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04-JUN-1998;
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17-JUN-1998
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Human, PRO, probe, ss, secreted polypeptide, transmembrane polypeptide, toxin, radiolabel, cell death, gene mapping, chromosome mapping, protein electrophoresis, genetic disorder, immunosuppressive, cytostatic,
hybridisation probes for mapping the gene which encodes the PRO polypeptide, and for the genetic analysis of individuals with genetic disorders, in gene therapy, for chromosome identification, as chromosome markers, and for generating probes for PCR, Northern analysis, Southern analysis and Western analysis. The present sequence represents a PCR primer used in the examples of the present
                                                                                                                                                                                      Gaps
                                                                                   Note: The sequence data for this patent was obtained in electronic format directly from the USPTO web site at seqdata.uspto.gov/psipsDIDEntry.html.
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                                                                                                                                                            Match 1.0%; Score 13.4; DB 1; Length 17; Local Similarity 93.3%; Pred. No. 2.1e+02; es 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                     Sequence 17 BP; 1 A; 4 C; 7 G; 5 T; 0 other;
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2000WO-US03565.
2000WO-US04341.
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99WO-US12252
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2000WO-US04914
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2000WO-US13358
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99WO-US28301
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                                                                                                                                                                                                                                                                                                   ABX17250 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                     04-FEB-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                            Human PRO probe #59.
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24-FEB-2000;
24-FEB-2000;
02-MAR-2000;
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06-JAN-2000;
11-FEB-2000;
18-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                          antibacterial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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16-DEC-1999;
20-DEC-1999;
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15-SEP-1999;
15-SEP-1999;
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08-MAR-1999
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01-DEC-1999
                                                                          invention.
                                                                                                                                                                                                                                                                                                                             ABX17250;
                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                            RESULT 357
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WPI; 2003-066810/06.

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970S-065211P
970S-065111P
980S-065111P
980S-078910P
980S-081600P
980S-081600P
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980S-081600P
980S-08102P
980S-08103P
980S-08810P
980S-08810P
980S-088142P
980S-088142P
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98US-089440P.
98US-089512P.
98US-089514P.
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2001WO-US21735.
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2000WO-US23328
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2000WO-US32678
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2001WO-US17800
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                                     2000WO-US22031
                            28-JUL-2000)
11-AUG-2000)
23-AUG-2000)
24-AUG-2000)
01-DEC-2000)
01-DEC-2001)
01-JUN-2001)
20-JUN-2001)
29-JUN-2001)
16-JUN-1991)
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16-JUN-1998;
16-JUN-1998;
17-JUN-1998;
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28-APR-1998;
07-MAY-1998;
28-MAY-1998;
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24-NOV-1997
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## (GETH ) GENENTECH INC.

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DL;
Godowski PJ;
Paoni NF;
Wood WI;
  Eaton
i AJ, Baker KP, Botstein D, Desnoyers L, Eaton N, Fong S, Gerber H, Gerritsen ME, Goddard A, JC, Gurney AL, Kljavin IJ, Napier MA, Pan J, Stewart TA, Tunas D, Watanabe CK, Williams PM,
                                  Grimaldi JC,
Roy MA, Stewa
Zhang Z;
  Ashkenazi AJ,
                        errara N,
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                                                                                                                                                                 The invention relates to a secreted and transmembrane polypeptide, termed PRO polypeptide, and the polynucleotide encoding it. The polypeptide is useful for detecting PRO polypeptides and for linking a bioactive molecule to a cell expressing the above polypeptides, where the bioactive causes the death of the cell. The polypeptide, The bioactive material agonists or antagonists of the PRO polypeptide, for preparing variants of PRO, as a molecular weight marker for protein electrophoresis purposes and the PRO polypucleotide is useful for recombinantly expressing those markers. The polynucleotide is useful for recombinantly expressing those markers. The polynucleotide is useful for recombinantly expressing those chromosome and gene mapping, in generation of antisense RNA and DNA, in the preparation of PRO polypeptide, for generating transgenic animals or knocked animals which in turn are useful in the development animals or knocked animals.
                                                                                                                                                                                                                                                                                                                                                                                                                                  screening of therapeutically useful reagents, to construct hybridisation probes for mapping the gene which encodes PRO and for the genetic analysis of individuals with genetic disorders, in gene therapy, for chromosome identification, as a chromosome marker and for generating probes for PCR, Northern analysis, Southern analysis and Western analysis. This sequence represents a human PRO probe of the invention.
                                                    Novel secreted and transmembrane polypeptide for modulating biological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunosuppressants; immunoenhancers; treatment; diagnosis; screening; immune disorders; transporter peptides; proteasome complex; MMC class I molecules; HLA; antigen processing; antigen presentation; autoimmune disease; ankylosing spondylitus; prenatal diagnosis; polymerase chain reaction; se
                                                                         activity of cell expressing the polypeptide, identifying agonists or antagonists of polypeptide, and as molecular weight markers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.1e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Trowsdale J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 1 A; 4 C; 7 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Control probe #4 for caucosoid RING11 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (IMCR ) IMPERIAL CANCER RES TECHNOLOGY.
                                                                                                                                      Example 170; Page 301; 655pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Glynne R, Kelly AP, Powis SH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ26549/c
ID AAQ26549 standard; DNA; 18 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93.38;
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91GB-0019711.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1992-250030/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
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16-SEP-1991;
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Claim 5; Fig 13; 287pp; English.
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94US-0354920.
94US-0363253.
94US-0363254.
95US-0390850.
95US-0426124.
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95US-0434509.
95US-0000951.
95US-0000974.
                                                                                                                                                                                                                                                                                           703 CICCITGATICIGIG 717
                                                                                                                                                                                                                                                                 93.3%;
92US-0987130
92US-0987133
                                                                                                                                                                                                                                                                                                                                                         AAX67098 standard; RNA; 18
                           (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                            creerraarrerare 1
                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 93.33
Matches 14, Conservative
                                            Dudycz LW,
                                                                       WPI; 1993-386599/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    diagnosis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUL-1995;
07-AUG-1995;
07-DEC-1992;
07-DEC-1992;
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                                            Draper KG,
Mamone JA;
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AAX67098
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                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                          RNA; enzyme; enzymatic RNA molecule; ERM; cleave; RNA; mRNA; HnRNA; picornavirus; HIV; immunodeficiency virus; hepatitis B virus; HBV; papilloma virus; HPV; Epstein-Barr virus; EBV; TCLV; TCLV; T-cell leukaemia virus; hepatitis C virus; HCV; cytomegalovirus; influenza virus; HSV; herpes simplex virus; vector; immune response;
                                                   This probe was used together with AAQ26546-51 to analyse caucosoid controls by oligonucleotide typing, whilst investigating RING 11 polymorphisms - see AAQ26544,5.
DNA encoding RING4, RING10, RING11 AND RING12 proteins - for treatment and diagnosis of immune disorders and screening of new immunosuppressants and immuno:enhancers
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0
                                                                                                         1.0%; Score 13.4; DB 1; Length 18; 93.3%; Pred. No. 2.3e+02;
                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                        antibody; ribozyme; viral RNA; treatment; ss.
                                                                                          Sequence 18 BP; 3 A; 6 C; 6 G; 3 T; 0 other;
                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                           Cytomegalovirus target sequence 18.
                                      Example 2; Page 40; 101pp; English
                                                                                                                                                                                                             AAQ52841 standard; RNA; 18 BP
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92US-0883823.
92US-0883849.
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92US-08828889.
92US-0882889.
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92US-0884333.
92US-0884422.
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92US-0987129
                                                                                                                                               625 GACCAGCTCCAGGAG 639
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92US-0882713
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                                                                                                                                                               GCCCAGCTCCAGGAG 2
                                                                                                                                                                                                                                                (updated)
(first entry)
                                                                                                                             14; Conservative
                                                                                                          Query Match
Best Local Similarity
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07-DEC-1992;
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14-MAY-1992;
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14-MAY-1992;
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14-MAY-1992;
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                                                                                                                                                                                                                                                         26-MAY-1994
                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                               AAQ52841;
                                                                                                                                                                16
                                                                                                                              Matches
                                                                                                                                                                                           RESULT 359
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Arthritic condition, graft tolerance, immune response, target; cleavage; hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                       The sequences (AAO52824-Q52880) are pref. Cytomegalovirus target sequences for enzymatic RNA molecules. The RNA molecules are complementary to a substrate binding region in the specified gene target. They also have enzymatic activity, in that they specifically cleave RNA in the target. The ERMs interfere with viral replication and therefore have anti-viral properties. They can be used to attenuate viruses to be used in vaccines.

(Updated on 25-MAR-2003 to correct PN field.)

(Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
Holecek JJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ò
                                                                                                                                                                                                 Enzymatic RNA molecules - used to inhibit viral replication, infection and gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.0%; Score 13.4; DB 1; Length 18; 13.3%; Pred. No. 2.3e+02; ve 0; Mismatches 1; Indels
    Macejak DG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human B7-2 hairpin ribozyme target SEQ ID NO:3730.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 8 A; 4 C; 4 G; 2 U; 0 other;
    Mcswiggen JA,
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(RIBO-) RIBOZYME PHARM INC.
  Draper K,
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Synovial membrane of joints for the treatment or prevention of arthritis, particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for treating graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysis without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly specific. The present sequence is used in the exemplification of the The present invention describes a novel enzymatic nucleic acid (ENA) having a hammerhead motif (HM) comprising: (i) at least 5 ribose residues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification. The ENA's can inhibit collagenase and stromelysin production in the Enzymatic nucleic acid molecules having a hammer-head motif - used for the treatment of arthritis, induction of graft tolerance or McSwiggen J, Pavco P, Stinchcon A, Modak A, Usman N, Burgin A; s T, Thompson JD, Wincott F; Sequence 18 BP; 3 A; 3 C; 2 G; 10 U; 0 other; Claim 10; Page 216; 307pp; English. treatment of auto-immune diseases 1.0%; Beigelman L, Karpeisky A, Matulic-Adamic J, Jarvis T, WPI; 1996-300653/30. Query Match Best Local Similarity

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Stinchcomb DT;

Gustofson J,

Score 13.4; DB 1; Length 18; Pred. No. 2.3e+02; 9; Mismatches 1; Indels 9; 1111 GTTTTCTGTTTAATT 1125 |::::|:|::|
GUUUUCUGUCUAAUU 15 Conservative Matches ð g

. 0

Gaps

. 0

Nucleotide sequence PCR primer 8. AAV54171 standard; cDNA; 18 BP. (first entry) 21-DEC-1998 AAV54171; RESULT 361 AAV54171 

PCR; primer; amplification; apoptosis; antibody; inhibition; immunohistological staining.

58

Synthetic.

W09839437-A1

11-SEP-1998

98WO-JP00905 97JP-0050302 05-MAR-1998; 05-MAR-1997;

(KYOW ) KYOWA HAKKO KOGYO KK.

Sakaki Y;

WPI; 1998-495844/42

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is an antisense oligonucleotide used to modulate expression of Galpha-16. Galpha-16 is a human G protein which interacts differentially with several receptor types including members of the opioid and chemokine receptor families. A series of antisense oligonucleotides have been designed to target different regions of human Galpha-16 RNA. They may be used to inhibit the expression of Galpha-16 in human calls and tissues and thus to treat diseases associated with Galpha-16, such as hyperproliferative disorders, especially cancer. Infections, inflammation or tumour formation can be prevented or delayed. The compounds can be used in research and diagnostics in sandwich and other assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; G-alpha-16; G protein; cytostatic; hyperproliferative disorder; cancer; inflammation; infection; antisense inhibition; ss.
                                                                                                   This is the nucleotide sequence of a PCR primer used in the method of the invention, involving the use of novel apoptosis-related DNAs and proteins. The inventions can be used as diagnostic reagents for apoptosis e.g. (monoclonal) antibodies for the protein, as a reagent in immunohistological staining, as apoptosis inhibitors. It can also be used for treatment of apoptosis-related diseases.
                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Note: The sequence has a phosphorothicate backbone and may be either an oligodeoxynucleotide or a chimeric oligonucleotide containing 2'-methoxyethyl (2'-MOE) wings and a deoxy gap. The ISIS number given above corresponds to the oligodeoxynucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A new antisense compound for inhibiting the expression of human G-alpha-16 and treating, preventing or delaying infections, inflammation or hyperproliferative disorders such as cancer -
                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                Length 18;
Novel apoptosis-related DNAs and proteins - for diagnosis, preventing or treating diseases associated with apoptosis
                                                                                                                                                                                                                                                                                                                                           1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human G-alpha-16 antisense oligonucleotide ISIS# 20818.
                                                                                                                                                                                                                                                                                              1.0%; Score 13.4; DB 1;
93.3%; Pred. No. 2.3e+02;
iive 0; Mismatches 1;
                                                                                                                                                                                                                                                     Sequence 18 BP; 0 A; 0 C; 3 G; 15 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 15; Page 72; 100pp; English.
                                                            Example 1; Page 49; 70pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                     1144 TTTTTTTTTTGG 1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-US19613.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98US-0205143.
                                                                                                                                                                                                                                                                                                                                                                                                                              rrrrrrrrrdg 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA48761 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                           Ouery Match
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-412354/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200032817-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SISI (-SISI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-JUN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA48761;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA48761/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      achieves higher signal to noise ratios to increase the detection limits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Signal processing method useful for achieving higher signal to noise ratios and to increase detection limits of target analytes
                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                      Detect; target analyte; electrode array; environmental pollutant; pesticide; insecticide; toxin; chemical; virus; spore; ss.
                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                   DNA sequence #4 used in target nucleic acid detection method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
1.0%; Score 13.4; DB 1; Length 18;
Best Local Similarity 93.3%; Pred. No. 2.3e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                     Length 18;
                                                                                      Indels
                                                Query Match
1.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 2.3e+02;
Matches 14; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 3 A; 2 C; 7 G; 6 T; 0 other;
               Sequence 18 BP; 4 A; 8 C; 2 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'note = "OTHER = T(CH2)16SH"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Page 122; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CLIN-) CLINICAL MICRO SENSORS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                          AAA10983 standard; DNA; 18 BP.
                                                                                                                           1336 GTGTTTCAGGCAGGG 1350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-US21683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0100730
                                                                                                                                                           16 Grerrreaggeaagg 2
                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of the target analytes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-271554/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200016089-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-SEP-1998;
                                                                                                                                                                                                                                                                                                                               20-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               O'Connor SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                            AAA10983;
                                                                                                                                                                                                                   RESULT 363
                                                                                                                                                                                                                                  AAA10983
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This sequence represents a signalling probe used in an example of the method of the invention. The invention relates to the detection of target analytes in a sample using an electrode array. At least one electrode forms an assay complex consisting of a capture binding ligand covalently attached to the electrode, a target analyte, and an electron transfer moiety. An input signal is applied to the assay complex, and the resulting output signal is processed to detect the presence of the target analytes. Preferred analytes include proteins and nucleic acids. However, the analyte may also be an environmental pollutent (pseticides, insecticides or toxins), a chemical (solvent or organic material), biomolecules (hormones, cytokines, proteins, lipids, carbohydrates, cellular membrane antigens and receptors), whole cells (prokaryotic and spores. The method achieves higher signal to noise ratios to increase the detection limits of the target analytes.
                                                                                                                                                                                                                                                             Detect, target analyte, electrode array, environmental pollutant; ss, pesticide, insecticide, toxin, chemical; virus, spore, signalling probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Signal processing method useful for achieving higher signal to noise ratios and to increase detection limits of target analytes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                            Partial signalling probe sequence used in analyte detection method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%; Score 13.4; DB 1; Length 1. 13.3%; Pred. No. 2.3e+02; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 6 A; 7 C; 2 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 122; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CLIN-) CLINICAL MICRO SENSORS INC.
                                                                                                                       BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93.3%;
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21
                                 4 GCAGTTGACGTGGAT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7 GCAGTTGAGGTGGAT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15 GCAGTTGACGTGGAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ90641 standard; DNA; 18
                                                                                                                     AAA10986 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-JUN-2000 (first entry)
                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 93.3
Matches 14; Conservative
 GCAGTTGAGGTGGAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-271554/23.
                                                                                                                                                                                                                                                                                                                                                  WO200016089-A2,
                                                                                                                                                                                                                                                                                                                                                                                                                      17-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-SEP-1998;
                                                                                                                                                                                         20-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              O'Connor SD;
                                                                                                                                                                                                                                                                                                                                                                                     23-MAR-2000
                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ90641;
                                                                                                                                                        AAA10986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 365
                                                                                    RESULT 364
                                                                                                      AAA10986,
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Gaps

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98JP-0225228.

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Human adipose tissue gene amplifying primer #2.
                                                                                                                                                                                                Sequence 18 BP; 0 A; 0 C; 3 G; 15 T; 0 other;
                                                                                                                                 Example 2; Page 18; 50pp; Japanese.
                                                                                                                                                                                                                                                                           AAZ97002 standard; DNA; 18 BP
                                                                                                                                                                                                                                    1144 TTTTTTTTTTGG 1158
                                                                                                                                                                                                                                                                                                              Nucleotide sequence of DNA4.
                                                                                                                                                                                                                                              4 rrrrrrrrrrrsd 18
                                                                                         (NISB ) JAPAN TOBACCO INC
                                                                                                     WPI; 2000-306578/27
                                                                                                                                                                                                                  Best Local Similarity
Matches 14; Conserv
                                            JP2000037190-A.
                                                                                                                                                                                                                                                                                                                                                                  Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                      23-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                               W09967425-A2
                                                                              23-JUL-1998;
                                                                                                                                                                                      cissue genes
                                 Homo sapiens
                                                                   23-JUL-1998;
                                                                                                                                                                                                                                                                                                  14-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                          29-DEC-1999
                                                       08-FEB-2000
                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                       AAZ97002;
                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                RESULT 366
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1.0%; Score 13.4; DB 1; 93.3%; Pred. No. 2.3e+02; ive 0; Mismatches 1;

Conservative

Location/Qualifiers

(first entry)

a "T (CH2) 16SH"

/\*tag= /note= 99WO-US14191

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detection electrode with a covalently attached capture ligand and at least comprises concentrating the analyte in a detection chamber comprising a detection electrode with a covalently attached capture ligand to form an cassay complex comprising the analyte, ligand and at least 1 electron transfer moiety (ETM), and detecting the ETM with the the detection electrode. The method is particularly used for assaying nucleic acids or correctly but more generally the target analyte is any compound for which a binding partner is available, e.g. pesticide or other pollutants; the therapeutic or illicit drugs; whole cells; viruses etc. Particularly the capture ligands. Particular applications are detecting genes different capture ligands. Particular applications are detecting genes of effecting bacteria and viruses (e.g. for blood screening or testing water or foods); for forensic fingerprinting; for sequencing and for detecting successful gene amplification. The rate of complex formation is concessed, resulting in more sensitive detection, particularly down to course of the invention for detection of target sequences.
                                                                                                                                                                                                                                                                                                              Complex-forming assay for analyte, particularly nucleic acid, used e.g. in detecting genetic disease, with accelerated complex formation by electrophoretic concentration of analyte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Target analyte; nucleic acid detection; capture ligand; pesticide; electron transfer moiety; pollutant; therapeutic drug; cancer; Alzheimer's disease; cystic fibrosis; blood screening; water testing; forensic fingerprinting; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention relates to a method of detecting a target analyte that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.0%; Score 13.4; DB 1; Length 18; 93.3%; Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleotide sequence of a 30nm signalling probe D-1055.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 3 A; 2 C; 7 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Page 113; 140pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CLIN-) CLINICAL MICRO SENSORS INC.
                                                                                                        (CLIN-) CLINICAL MICRO SENSORS INC.
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98US-0090389.
98US-0134058.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7 GCAGTTGAGGTGGAT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GCAGTTGACGTGGAT 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                               Ö
                                                                                                                                                                               Blackburn
                                                                                                                                                                                                                                                  WPI; 2000-136990/12.
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   23-JUN-1998;
                                         14-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-DEC-1999.
                                                                                                                                                                                   Kayyem JF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ97005;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 367
AAZ97005/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to identification of genes and proteins of adipose tissue relating to obesity, particularly complications of visceral obesity including diabetes, hyperlipemia, hypertension, arteriosclerosis, hyperuricemia and sleep apnea syndrome. The genes arteriosclerosis and the proteins (AAV67598 *V6760) are used in the genetic diagnosis, prevention and treatment of adipose tissue related diseases. Sequences AAZ90640-51 represent PCR primers amplifying the human adipose
                                                                                                        Adipose tissue; obesity; diabetes; hyperlipemia; hypertension; human; arteriosclerosis; hyperuricemia; sleep apnea syndrome; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Target analyte; nucleic acid detection; capture ligand; pesticide; electron transfer moiety; pollucant; therapeutic drug; cancer; Alzheimer's disease; cystic fibrosis; blood screening; water testing; forensic fingerprinting; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   physiologically active protein specifically derived from mammal
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0

O'connor SD;

Blackburn G,

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The invention relates to a method of detecting a target analyte that comprises concentrating the analyte in a detection chamber comprising a detection electrode with a covalently attached capture ligand to form an assay complex comprising the analyte, ligand and at least I electron transfer moiety (ETM), and detecting the ETM with the the detection transfer moiety (ETM), and detecting the ETM with the the detection transfer moiety (ETM), and detecting the ETM with the the detection transfer moiety (ETM), and detecting to ETM with the the detection transfer moiety (ETM), and detecting to ETM with the the detection of proteins, but more generally the target analyte is any compound for which a binding partner is available, e.g. pesticide or other pollutants, the method is used in array formats, optionally with many thousands of different capture ligands. Particular applications are detecting genes associated with cancer. Alzheimer's disease, cystic fibrosis etc.; detecting bacteria and viruses (e.g. for blood screening or testing water or foods), for forensic fingerprinting, for sequencing and for detecting successful gene amplification. The rate of complex formation is increased, resulting in more sensitive detection, particularly down to locable the present sequence represents a signalling probe used in the course of the invention for detection of target sequences.
                                                                                                              Complex-forming assay for analyte, particularly nucleic acid, used in detecting genetic disease, with accelerated complex formation by electrophoretic concentration of analyte
                                                                                                                                                                                                                                  Example 2; Page 113; 140pp; English
                                                    WPI; 2000-136990/12.
Kayyem JF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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. 0 Length 18; 1; Indels Score 13.4; DB 1; Pred. No. 2.3e+02; 0; Mismatches 1; Sequence 18 BP; 6 A; 7 C; 2 G; 3 T; 0 other; 1.0%; 7 GCAGTTGAGGTGGAT 21 14; Conservative Local Similarity

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Gaps

GCAGTTGACGTGGAT 1 15

AAZ44140 standard; DNA; 18 BP 24-MAR-2000 (first entry) RESULT 368 

Human EGR-1 DNA antisense primer #24162.

BGR-1; early growth response 1; antisense; inhibition; human; primer; anti-inflammatory; cytostatic; antiviral; detection; diagnosis; viral infection; inflammation; tumor; ss.

Homo sapiens

US6008048-A.

28-DEC-1999.

98US-0205921. 04-DEC-1998;

98US-0205921 04-DEC-1998;

(ISIS-) ISIS PHARM INC.

Cowsert LM;

Monia BP,

WPI; 2000-096375/08.

Antisense oligonucleotides that inhibit expression of human early growth response-1, useful for diagnosis, treatment and prevention of tumors, inflammation and infection

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This invention describes novel antisense oligonucleotides (I) capable of inhibiting expression of human EGR-1 (early growth response-1). The products of the invention have anti-inflammatory, cytostatic and antiviral activity. (I) was tested for its effects on EGR-1 mRNA levels by real-time polymerase chain reaction (PCR), results indicated that 60% inhibition was achieved. When (I) was modified by 2'-O-methoxyethyl substitution of the first 4 and last 4 residues, and by replacing any C in these flanking regions with 5-methyl-C, the degree of inhibition was increased to 71%. (I) is used to inhibit expression of EGR-1 in cells and tissues in vitro, for research or diagnosis, e.g. detecting EGR-1 encoding nucleic acid. (I) may also be used to treat or prevent EGR-1-associated diseases, particularly viral infections, inflammation and tumors. AAZ44169 represent antisense primers used to inhibit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense compound which modulates human sentrin expression, useful for treating diseases associated with sentrin expression -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes an antisense compound (I) 8-30 nucleotides long targeted to a nucleic acid molecule encoding human sentrin. The antisense compound comprises a phosphorothicate antisense oligonucleotide which inhibits expression of human sentrin. (I) is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human sentrin phosphorothioate antisense oligonucleotide SEQ ID NO:47.
                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; sentrin, antisense oligonucleotide; phosphorothioate;
inhibition; modulation; expression; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                             1.0%; Score 13.4; DB 1; Length 18; 33.3%; Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/note= "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 4 A; 8 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
              Claim 1; Column 37-38; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 15; Column 38; 29pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ35905 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                348 CAGTGGCCGAGTGAG 362
                                                                                                                                                                                                                                                                                                                                                             93.3%;
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                                                                                                                                                                                                                                                                                                                                                             Local Similarity 93.3
nes 14; Conservative
                                                                                                                                                                                                                                                                        the human EGR-1 protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-022284/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Baker BF,
                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ35905;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 369
                                                                                                                                                                                                                                                                                                                                                                                 Matches
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useful for inhibiting expression of sentrin in human cells or tissues in vitro, for treating humans or other animals suspected of having or being prone to a disease associated with sentrin expression. (I) can also be used for research or diagnostic purposes. The present sequence represents a human sentrin phosphorothioate antisense
                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                               .
                                                                                                                                                                                                                Length 18;
                                                                                                                                                                                                            1.0%; Score 13.4; DB 1; Length 1 93.3%; Pred. No. 2.38+02; cive 0; Mismatches 1; Indels
                                                                                                                                                                  Sequence 18 BP; 4 A; 5 C; 3 G; 6 T; 0 other;
                                                                                                                 oligonucleotide from the present invention.
                                                                                                                                                                                                                                                             14; Conservative
                                                                                                                                                                                                                                    Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                Query Match
  8888888888888
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440 GAAAGTTGCTGAAGT 454 GAAAGTTACTGAAGT 4 18 ð d

AAH75239 standard; DNA; 18 02-OCT-2001 AAH75239; RESULT 370 AAH75239 

BP.

(first entry)

Human inducible NOS antisense oligonucleotide SEQ ID NO 83.

Antisense oligonucleotide; inducible nitric oxide synthase; NOS; modulate expression; immunomodulator; antidiabetic; cardiovascular; cardiant; neuroprotective; vasotropic; ischaemia; reperfusion injury; 2'-0-methoxyethyl; phosphorothioate; human; ss.

Homo sapiens,

ocation/Qualifiers 1..18 Key modified\_base

'note= "phosphorothioate backbone, 5' and 3' four nucleotide 2'-MOE (2'-0-methoxyethyl) wings, all cytidine residues are 5-methylcytidines and a deoxy gap" mod base= OTHER ಗ \*tag=

WO200152902-A1

26-JUL-2001.

15-JAN-2001; 2001WO-US01381

24-JAN-2000; 2000US-0490208

(ISIS-) ISIS PHARM INC

Bennett CF, Dean NM,

Cowsert LM;

WPI; 2001-465340/50.

New antisense oligonucleotides for modulating the expression of inducible nitric oxide synthase in cells or tissues, particularly useful for treating e.g. immunological, cardiovascular or neurological disorders, or ischaemia

Claim 3; Page 84; 144pp; English.

The invention relates to antisense compounds, especially oligomucleotides, which are targeted to a nucleic acid encoding inducible nitric oxide synthase and which specifically hybridise to and modulate expression of inducible nitric oxide synthase. The antisense compounds have immunomodulator, antidiabetic, cardiovascular, cardiant, neuroprotective, disorder and vasotropic activity. The antisense oligomucleotides are useful for inhibiting the expression of inducible nitric oxide synthase in cells or tissues. In particular, the antisense

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oligonucleotides are useful for treating diseases or disorders associated
                      with inducible nitric oxide synthase, e.g. diabetes, immunological disorder, cardiovascular disorder, neurological disorder or ischaemia-reperfusion injury. The antisense oligonucleotides are also useful for research and diagnostics. The present sequence is that of an antisense 2'-O-methoxyethyl gapmer oligonucleotide with a phosphorothicate backbone, a central "gap" region of ten nucleotides flanked by four nucleotide 2'-MOE (2'-methoxyethyl) wings and 5-methylcytidine residues throughout the oligonucleotide. The antisense olignucleotide is targeted to human inducible nitric oxide synthase (NOS)
                                                                                                                                                                                                                                                                                                         mRNA (AAH47973)
222222222222222233
2222222222222233
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Sequence 18 BP; 5 A; 6 C; 6 G; 1 T; 0 other;

Gaps ö Score 13.4; DB 1; Length 18; Pred. No. 2.3e+02; 0; Mismatches 1; Indels 1.0%; Query Match
Best Local Similarity 93.3
Matches 14; Conservative

ö

552 GGCAGGCATGCACAC 566 4 GGCAGGCACGCACAC 18 ò qq

RESULT 371 **AAH558**81

BP. AAH55881 standard; DNA; 18

AAH55881;

(first entry) 04-SEP-2001 Human SCN1A PCR-SSCP PCR primer SEQ ID NO:125.

Human, epilepsy, chromosome 2; SCNIA; SCNIA; sCNIA; identification; diagnosis; mutation; chromosome 2q23-q31; neurological disorder; anticonvulsant; neuroprotective; PCR primer; ss.

Homo gapiens. Synthetic.

WO200138564-A2.

31-MAY-2001

24-NOV-2000; 2000WO-CA01404.

99US-0167623. 26-NOV-1999;

(UYMC-) UNIV MCGILL.

Cossette P, Rochefort D, LaFreniere RG, WPI; 2001-355945/37. Roulean GA, 

Ragsdale D;

Determining a predisposition to epilepsy and/or development of epilepsy comprises determining the genotype of SCNIA, SCNIA and/or SCNIA, or a DNA variant, equivalent, or mutation which shows a linkage disequilibrium -

Example 3; Fig 2; 268pp; English.

The present invention describes a method (M1) of determining an individual's predisposition to epilepsy and/or development of epilepsy, as well as predisposition to epilepsy and/or development of epilepsy, method comprises determining the genotype of at least one gene selected from SCN1A, SCN2A or SCN3A, or a DNA variant, equivalent, or mutation which shows a linkage disequilibrium. SCN1A, SCN2A and SCN3A are all sodium channel genes located on chromosome 2. The idiopathic generalised epilepsy (1GB) gene is more specifically localised on chromosome 2q23-q31. Compounds identified as modulators of the biological activity of SCN1A, SCN2A or SCN3A proteins or genes, are useful for treating epilepsy or other neurological disorders. They have anticonvulsant and neuroprotective activities. AAHS5763 to AAHS6164 and AAB99674 to

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AAB99679 represent SCN1A, SCN2A, and SCN3A oDNAs, gene fragments, PCR primers, oligonucleotides and proteins given in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The specification describes antisense oligonucleotides which are targeted to a nucleic acid encoding Her-4 (a receptor/tyrosine kinase). The antisense oligonucleotides are used to inhibit the expression of Her-4 in cells or tissues in vitro. They can be used in diagnostics, therapeutics, prophylaxis and as a probe in research reagents. The
                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Compound for inhibiting the expression of Her-4 (a receptor/tyrosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                             "all cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          kinase) e.g. in preventing tumour formation, comprises an antisense oligonucleotide that hybridizes to a nucleic acid encoding Her-4 -
                                                                                                                                                                                                                                                                                                                     Antisense oligonucleotide; Her-4; receptor kinase; tyrosine kinase; infection; inflammation; tumour; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                            Antisense oligonucleotide targeted to human Her-4 coding region.
                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                             a
"2'-methoxyethyl (2'-MOE) nucleotides"
                                                                              Length 18;
                                                                                                      1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'note= "all internucleoside linkages are
  phosphorothioate linkages"
                                                                           Score 13.4; DB 1;
Pred. No. 2.3e+02;
0; Mismatches 1;
                                                   Sequence 18 BP; 4 A; 6 C; 6 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "2'-deoxynucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Column 43-44; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                      0;
                                                                                                                              1082
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                                                                             1.0%;
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                                                                                                                                                                                                                 AAH25339 standard; DNA; 18
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                                                                                                    Conservative
                                                                                                                           1068 CATCAGGCAGGCTCT
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/*tag=
/note=
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15..18
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                          the present invention.
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                                                                         Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                             modified base
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                                                                                                                                                                                                                                                                    22-AUG-2001
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                                                                                                                                                                                          RESULT
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antisense oligonucleotides can be used to prevent or delay infection, inflammation or tumour formation. AAH25315-AAH25398 represent antisense oligonucleotides which are targeted to different regions of the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Substrate composition, used of target analytes such as biomolecules, comprises a surface having an array of detection electrodes, each with a covalently attached capture ligand -
                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                               DNA array; DNA binding acceleration; analyte detection; sequencing; genetic diagnosis; pathogen detection; DNA fingerprinting; probe; gene probe assay; ss.
                                                                                                                                         0
                                                                                                            Length 18;
                                                                                                         1.0%; Score 13.4; DB 1; Length 1
93.3%; Pred. No. 2.3e+02;
tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CH2)16SH"
                                                                             6 C; 6 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 3 A; 2 C; 7 G; 6 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= "OTHER"
/note= "modified by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 113; 146pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CLIN-) CLINICAL MICRO SENSORS INC.
                                                                                                                                                                                                                                                                            BP.
                                                                                                                                                                                                                                                                                                                                                                    DNA array oligonucleotide #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Vielmetter JG,
                                                                                                                                                                     GGCAGGCATGCACAC 566
                                                                                                                                                                                                   GGCAGGCACGCACAC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-NOV-2000; 2000WO-US31233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0440371
                                                                                                                                                                                                                                                                          AAH25861 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                       14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag=
                                                                             BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-389777/41.
                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200135100-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-NOV-1999;
23-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Blackburn G,
                                                                                                                                                                                                                                                                                                                                      22-AUG-2001
                                                                             Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-MAY-2001
                                               Her-4 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                        AAH25861;
                                                                                                                                                                     552
                                                                                                         Query Match
Best Local 8
                                                                                                                                                                                                                                              RESULT 373
                                                                                                                         Best Loca
Matches
                                                                                                                                                                                                                                                            AAH25861
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1.0%; Score 13.4; DB 1; Length 18;

Query Match

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/note= "(C23)4-N87-N87-N87-N87-adenosine, where each N87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a composition comprising two nucleic acids each containing an electron-transfer group (ETM) having different redox potentials. The invention is used for electronic detection of nucleic acids, especially of substitutions (mismatches) and single-nucleotide polymorphisms, e.g. for genotypin and monitoring gene expression. The present sequence is a probe used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acids containing electron-transfer group, useful as labels in hybridization assays, e.g. for genotyping, allowing repeat analyses on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence determination using electronic detection signalling probe #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NOTE: Revised record submitted with a corrected sequence)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
Electron-transfer group; ETM; mismatch; genotyping;
gene expression; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Electron-transfer group; ETM; mismatch; genotyping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.0%; Score 13.4; DB 1;
93.3%; Pred. No. 2.3e+02;
iive 0; Mismatches 1;
                                                                                                                 /*tag= a
/mod_base= OTHER
/note= "modified by (CH2)16SH"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 3 A; 2 C; 7 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Page 111; 159pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                     Location/Qualifiers
18
                                                                                                                                                                                                                                                                                                                                    (CLIN-) CLINICAL MICRO SENSORS INC.
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF58216 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                 26-JUL-2000; 2000WO-US20476.
                                                                                                                                                                                                                                                                                26-JUL-1999; 99US-0145695
17-MAR-2000; 2000US-0190259.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7 GCAGTTGAGGTGGAT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene expression; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 GCAGTTGACGTGGAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (updated)
                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-159728/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    a single surface
                                                                                                                                                                                  WO200107665-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
modified_base
                                                                                Key
modified_base
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24-APR-2001
                                                                                                                                                                                                                 01-FEB-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF58216;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                      Umek RM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF58216/
 g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention provides the protein and coding sequences of leptin fragments which modulate the activity of lipolysis stimulated factor (LSR). These sequences are useful in the treatment of obesity related diseases, including obesity, anorexia, carchexia, cardiac and coronary insufficiency, stroke, hypertension, atheromatous disease, atherosclerosis, non-insulin dependent diabetes, hyperlipidaemia,
               ·,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New leptin polypeptide fragment and related polynucleotides, useful for
the prevention and treatment of obesity and obesity-related diseases
such as hypertension and diabetes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                 Gaps
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0
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                                                                                                                                                                                                                                                                                                Leptin, human; LSR; lipolysis stimulated receptor; obesity; hypertension; anorexia; cachexia; stroke; atherosclerosis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.0%; Score 13.4; DB 1; Length 18; 93.3%; Pred. No. 2.3e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence determination using electronic detection probe #4.
               Indels
                                                                                                                                                                                                                                                                  Zinc finger coding sequence related oligo SEQ ID NO: 91.
93.3%; Pred. No. 2.3e+02;
ive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 2 A; 2 C; 12 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bihain B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 12; Page 244; 247pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fruebis J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF58213 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hyperuricaemia and syndrome X.
                                                                                                                                                                AAF62366 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          186 CCCCGCCGCCCACCC 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-SEP-2000; 2000WO-IB01470
                                                                  4 GCAGTTGACGTGGAT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0155506
                                                  21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (updated)
(first entry)
                                                                                                                                                                                                                                 (first entry)
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Matches 14, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18 CCCCGCCGCTCACCC
               14; Conservative
                                               7 GCAGTTGAGGTGGAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yen F, Erickson MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-218642/22
 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                    WO200121647-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-SEP-1999;
                                                                                                                                                                                                                               06-JUN-2001
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24-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                   29-MAR-2001
                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF58213;
                                                                                                                                                                                                 AAF62366;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 375
                                                                                                                                 RESULT 374
               Matches
                                                                                                                                                AAF62366/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF58213
ID AAF
XX
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olek A,
Lewin A,
Pelet C,
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                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 378
                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ7554
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                                                                                                                                                                                                                                       The present invention relates to a composition comprising two nucleic acids each containing an electron-transfer group (ETM) having different redox potentials. The invention is used for electronic detection of nucleic acids, especially of substitutions (mismatches) and single-nucleotide polymorphisms, e.g. for genotypin and monitoring gene expression. The present sequence is a probe used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Haematopoietic cell proliferation disorder related oligonucleotide #982.
group has (C23)4 appended, where N87 is a branch point comprising a ring structure and C23 is shown in Fig IF PCTUSS99/01705"
                                                                                                                                                                               Nucleic acids containing electron-transfer group, useful as labels in hybridization assays, e.g. for genotyping, allowing repeat analyses on
                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; haematopoietic cell proliferation disorder; cytostatic; gene therapy; lymphocytic leukaemia; acute myelogenous leukaemia; cytosine methylation state; probe; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mueller J;
                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                               1.0%; Score 13.4; DB 1; Length 18; 3.3%; Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                          submitted with a corrected seguence)
                                                                                                                                                                                                                                                                                                                                                                  i, Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Braun A, Distler J, Guetig D, Howe A,
                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 6 A; 7 C; 2 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                     Example 2; Page 111; 159pp; English.
                                                                                                                        (CLIN-) CLINICAL MICRO SENSORS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                        93.38;
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                                                                          26-JUL-2000; 2000WO-US20476.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-2002; 2002WO-EP03401
                                                                                            26-JUL-1999; 99US-0145695
17-MAR-2000; 2000US-0190259
                                                                                                                                                                                                                                                                                                                                                                                    7 GCAGTIGAGGIGGAT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABZ10842 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                     15 GCAGTTGACGTGGAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (EPIG-) EPIGENOMICS AG.
                                                                                                                                                            WPI; 2001-159728/16.
                                                                                                                                                                                                   single surface
                                                                                                                                                                                                                                                                                                           (NOTE: Revised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200277272-A2
                                    WO200107665-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-0CT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-JAN-2003
                                                        01-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Berlin K,
                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABZ10842;
                                                                                                                                          Umek RM;
                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 377
                                                                                                                                                                                                                                                                                                                                                                                                                                            ABZ10842/
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differentiating between hematropoietic cell proliferative disorders associated with at least 1 gene and/or their regulatory regions in a subject. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least 1 reagent, which distinguishes between methylated and non-methylated CpG dinuclectides within the target nucleic acid. ABZ0961 to ABZ11118 represent specifically claimed nucleotide sequences from the present invention. Oligonuclectides from the present invention can be used: for differentiating between healthy haematopoietic cells and proliferative dissorder haematened from the present invention can be used: for differentiating between healthy haematopoietic cells and proliferative determining the cytosine methylation state and/or single nucleotide polymorphisms (SNPs) of haematopoietic cell proliferation disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        related sequences and their complements; and as primers for the amplification of haematopoletic cell profiferation disorder related DNA sequences. The nucleotide sequences from the present invention can also be used for detecting a predisposition to, differentiation between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               subclasses, diagnosis, prognosis, treatment and/or monitoring of haematopoietic cell proliferative disorders. The present method enables a highly specific classification of haematopoietic cell proliferative disorders allowing for improved and informed treatment of patients.
                                                                                                                                                                                                           Detecting and differentiating between hematopoietic cell proliferative disorders, comprises contacting a target nucleic acid with a reagent that distinguishes between methylated and non-methylated CpG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
I, Adorjan P, Grabs G, Lesche R, Leu E;
Maier S, Model F, Mueller V, Otto T;
Ziebarth H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a method for detecting and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 13.4; DB 1;
Pred. No. 2.3e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 6 A; 0 C; 7 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                               Claim 15; Page 65; 117pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              54 TACTCCTCAATTACC 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93JP-0112515
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17 racrecreatriate 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 93.3
es 14; Conservative
Piepenbrock C,
Lipscher E, M
                                 Lipscher B,
Schwope I,
                                                                                                                                       WPI; 2003-018942/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1995-018287/03
                                                                                                                                                                                                                                                                                                                        dinucleotides
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESE) files AAQ75547-Q7598) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression
                 cDNA and gene expression - by amplification of mRNA digestion with restriction enzymes
                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 0 A; 0 C; 2 G; 17 T; 0 other;
                                                                                          Disclosure; Page 5; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                          rapidly and easily.
                   Analysis of
followed by
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1.0%; Score 13.4; DB 1; Length 19; 93.3%; Pred. No. 2.4e+02; 1; Indels Mismatches 0 1144 TITITITITIGG 1158 Query Match
Best Local Similarity 93.3
Matches 14; Conservative

TTTTTTTTTTTTGG 19 Ŋ

AAX06004 standard; DNA; 19 BP AAX06004; RESULT 379 

10-MAY-1999 (first entry)

Oligo used in construction of plasmid pGP2

gene; Human; Immunoglobulin transgene; Ig; VH gene; D gene; JH gene; mu switch sequence; gamma gene; IgM; IgG; ss.

Synthetic.

US5874299-A

23-FEB-1999

97US-0800353 14-FEB-1997;

92US-0834539. 90US-0574748. 90US-0575962. 91WO-US06185 97US-0800353 31-AUG-1990; 29-AUG-1990; 14-FEB-1997 28-AUG-1991

(GENP-) GENPHARM INT INC.

Kay RM, Lonberg N;

WPI; 1999-179989/15.

Human immunoglobulin transgene - with mu and gamma isotype switching Example 5; Column 33; 88pp; English segments

The invention relates to a heavy chain (human) immunoglobulin (Ig) transgene. The transgene comprises: (1) human VH gene segments; (ii) human D gene segments; and either (iv) a mu constant region comprising a mu switch sequence upstream from a mu coding segment; (v) a gamma constant region comprising a gamma switch acquence upstream from a human gamma coding segment; where (vi) the mu and gamma constant regions are closer than in wild type human Ig heavy chain loci; or (vii) a heavy chain enhancer; (viii) a mu constant region

comprising a mu switch sequence upstream from a mu coding segment; (ix) a gamma constant region comprising a gamma switch sequence upstream from a human gamma coding segment; and (x) at least one discontinuity of at least 2 kb between the mu and gamma gene segments as compared to a human germ-line heavy chain locus; or (xi) a human mu CH gene and at least two non-mu human CH genes and their associated isotype switching sequences; where (xii) the human mu and human gamma switch sequences are closer than in wild type human ig heavy chain loci. The transgenes allow non-human animals to produce heterotogous (human) ig's with varying specificities. The presence of mu and gamma switch segments allows isotype switching of the human heavy chain mini-locus from IgM (for maturation) to IgG. 

Seguence 19 BP; 3 A; 7 C; 7 G; 2 T; 0 other;

ö Gaps ; Length 19; Indels 1.0%; Score 13.4; DB 1; 13.3%; Pred. No. 2.4e+02; ve. 0; Mismatches 1; 93.38; Local Similarity 93.3 nes 14; Conservative Query Match Matches

391 GTGGCAGCAATGGCC 405 GTGGCCGCAATGGCC 18 4 à d

ö

Gaps

.; 0

RESULT 380 AAS45588

BP. AAS45588 standard; DNA; 19

AAS45588; 

Human PARP-2 RT-PCR reverse primer.

(first entry)

18-DEC-2001

PARP; Poly (ADP-ribose) polymerase; antisense oligonucleotide; cytostatic; nootropic; neuroprofective; antiinflammatory; antidiabetic; immunosuppressant; hyperproliferative disorder; cancer; cellular injury; oxidative stress; neurological disorder; parkinsonism; apoptosis; mencingitis-associated intracranial complication; ischaemia; PCR primer; inflammatory disorder; autoimmune disorder; arthritis; diabetes.

Homo sapiens

WO200164955-A1

07-SEP-2001.

01-MAR-2001; 2001WO-US06572.

02-MAR-2000; 2000US-0517467.

(ISIS-) ISIS PHARM INC.

Popoff I,

Ë Cowsert

WPI; 2001-602570/68.

Antisense compound useful for treating hyperproliferative, neurological, inflammatory and autoimmune disorders and diabetes inhibits human PARP —

Example 13; Page 80; 168pp; English.

The invention relates to antisense oligonucleotides targeted to human PARP mucleic acid and inhibiting expression of human PARP. PARP PARP mucleic acid and inhibiting expression of human PARP. PARP PARP relations of partial partia

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0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Collection of binding groups for determining or typing samples, especially clinical samples, has groups capable to identify essentially all members of the family of nucleic acids of relatively high
                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                           HIV-1 related binding molecule oligonucleotide sequence SEQ ID NO:97.
arthritis) and diabetes. The present sequence is an RT-PCR (reverse transcriptase PCR) primer used to quantitate PARP mRNA levels.
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                                                                                          Length 19;
                                                                                                                               Indels
                                                                                        / Match 1.0%; Score 13.4; DB 1; Local Similarity 93.3%; Pred. No. 2.4e+02; nes 14; Conservative 0; Mismatches 1;
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                                                      Sequence 19 BP; 4 A; 9 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (AMST-) AMSTERDAM SUPPORT DIAGNOSTICS BV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Goudsmit J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus type 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure, Page 30; 166pp; English.
                                                                                                                                                                                                                                                                                               ABL88875 standard; DNA; 19 BP.
                                                                                                                                                                  229 CAGCCTCAGGCATCT 243
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-JUL-2000; 2000EP-0202611
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                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     significance
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                                                                                          Query Match
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DP
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                                                                                                                                                                      Probe to polymorphic site of the second exon of DPBeta allele of HLA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Target region for probe is amplified by polymerase chain reaction, probe binds to complementary regions in very stringent conditions, identifying variations in the hypervariable DP gene. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymerase chain reaction; PCR; amplify; primer; probe; HLA-DP; genotype; sequence specific; polymorphic region; variable segment; panel; autoimmune disease; insulin-dependant diabetes mellitus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIA DP genotyping by amplifying target DNA then hybridisation with panel of sequence specific oligonucleotide(s), esp. for assessing risk of autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                        HLA; autoimmune diseases; probe; DP gene; DPbeta allele.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.0%; Score 13.2; DB 1;
83.3%; Pred. No. 2.5e+02;
cive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 5 A; 5 C; 6 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HLA-DP genotype determination probe #26.
                                                                                                                                                                                                                                                                                                                                                                                                         Bugawan TL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 8; Page 48; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     521 ACCTGCCGGAGGAGCAGC 538
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 1144 TTTTTTTTTTTGG 1158
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88US-0258212.
89US-0347506.
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           AAN92661 standard; DNA; 18
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                                                                                                                                  (updated)
(first entry)
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1989-370738/50.
                                                                                                                                                                                                                                                                                                                                                                                 CORP.
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                                                                                                                                                                                                                                                                                                                                                                                (CETU ) CETUS
                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                        18-MAY-1989;
                                                                                                                                                                                                                                                                                                                                            14-OCT-1988;
04-MAY-1989;
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29-JUN-1994
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16-MAY-1990
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                                                                                                                                                                                                                                                                                30-NOV-1989
                                                                                                         AAN92661;
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AAQ54537
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Gaps

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1; Indels

Query Match
1.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 1;

Length 19;

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AAQ89234-Q89253 are PCR primers for fragments of hepatits C virus (HCV) or non-A non-B hepatits virus (NANBH) cDNA. These cDNAs code for amino acid sequences, which are antigens to structural and non-structural regions of the HCV virus. These antigens can be used in the diagnosis of NaNBH patients and the detection of HCV carriers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vascular endothelial growth factor receptor; VBGF receptor; flt-1; flt. TOR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; coular disease; fms-like; tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes nucleic acid molecules which modulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                    1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 2.5e+02;
                                                                                                                                                   A nucleic acid fragment coding Non-A Non-B Hepatitis virus antigens - for diagnosis of NANBH and detection of HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse flt-1 VEGF receptor hairpin ribozyme substrate #102.
                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                        ZH TOKYOTO RINSHO IGAKU SOGO KENKYUSHO.
                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 12 A; 2 C; 2 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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                                                             8
                                                                                                                                                                                                Example 1; Page 7; 22pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; Page 189; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Escobedo J, McSwiggen J, Pavco
                                                                                                                                                                                                                                                                                                                                                                                                                 1139 ATGCCTTTTTTTTTT 1156
                                                             SANWA KAGAKU KENKYUSHO
TONEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
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95US-0005974.
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                92JP-0207391
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RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX75643 standard; RNA; 18
                                              KOKUSAI SHIYAKU KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             foetal liver kinase 1; ss.
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                      WPI; 1994-298800/37.
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26-OCT-1995;
              10-JUL-1992;
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(TOFU) T
(TOKR-)
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(RIBO-)
                                              (KOKU-)
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                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                individuals susceptibility to an autoimmune disease including individuals susceptibility to an autoimmune disease including insulin-dependant diabetes mellitus and coeliac disease (CD) by disturbing the HIA-DP genotype. CD is linked to the alleles DPB13, DPB1, DPB1, DPB3 or DEB4.2. Susceptibility to particular juvenile rheumatoid arthritis is indicated by the DPB2.1 allele. The process can also be used to provide forensic evidence.

(Updated on 25-WAR-2003 to correct P1 field.)
                                                                                                                                                                                                                                                                                                                                                               The sequences given in AAG54509-614 are primers and probes which were used in the method of the invention for determining the HLA-DP genotype of an individual. The method comprises amplifying a target region of the nucleic acid in the sample in question, under conditions suitable for carrying out PCR with a sequence specific primer. The target region contains a polymorphic region (variable segment) of an HLA DP gane. The amplified sequences are mixed with a panel of probes, where each probe is complementary to a varient sequence of a variable segment of an HLA DP gene, than the DP gene, under stringent binding conditions. The method is useful for determining an
                                                                                                                                                                                                                                                                                     Method for HLA - DP typing - comprises amplfication of target nucleic acid region and hybridisation with oligo-nucleotide probes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Non-A non-B hepatitis virus antigens; NANBH; hepatitis C virus; ss.
coeliac disease; CD; allele; DPB13; DPB1; DPB3; DPB4.2; juvenile; rheumatoid arthritis; DPB2.1; forensic evidence; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 13.2; DB 1;
Pred. No. 2.5e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis C virus 8003-9388 fragment PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 5 A; 5 C; 6 G; 2 T; 0 other;
                                                                                                                                                                                                                             Erlich HA;
                                                                                                                                                                                                (HOFF ) HOFFMANN LA ROCHE & CO AG F.
                                                                                                                                                                                                                                                                                                                                      Claim 2; Page 81; 85pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       538
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       521 ACCTGCCGGAGGAGCAGC
                                                                                                                                     93EP-0109447
                                                                                                                                                                   92US-0903028
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                                                                                                                                                                                                                             Begovich AB, Bugawan TL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                            WPI; 1994-001049/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP06225770-A
                                                                                                                                     14-JUN-1993;
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                                                                        EP575845-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15;
                                                                                                      29-DEC-1993
                                            Synthetic.
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Matches
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AAQ89247/c
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or

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Gaps

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Length 18;

Query Match

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the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fims-like tyzosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and theumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAK67275 to AAX75752 represent specific examples
                                                                                                                                                                                                                                                                                                                                                                                                                                          Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KOR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene expression or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                           Length 18;
                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                 Human KDR VEGF receptor hairpin ribozyme substrate #2.
                                                                                                             of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid molecule modulating VEGF receptor(s)
                                                                                                                                                                       Score 13.2; DB 1;
Pred. No. 2.5e+02;
4; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stinchcomb
                                                                                                                                           Sequence 18 BP; 3 A; 7 C; 4 G; 4 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               McSwiggen J, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; Page 118; 218pp; English.
                                                                                                                                                                                                                              1053 CAGCCCTGGCCTTCCCAT 1070
                                                                                                                                                                                                                                              |||| ||:| ||:| ||:
CAGGCCUGACCUUCGCAU 18
                                                                                                                                                                     1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96US-0584040
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                              AAX71704 standard; RNA; 18
                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                   11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-259017/23.
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                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                     28-JUL-1999
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                                                                                                                                                                     Query Match
Best Local S:
Matches 11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vaccular endothelial growth factor (VEGF). A patient preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                      Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid molecule modulating VEGF receptor(s) gene expression or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mRNA stability - useful for treating e.g. tumour angioge
psoriasis, rheumatoid arthritis, etc., in a human patient
                                 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                           Human flt1 VEGF receptor hairpin ribozyme substrate #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Stinchcomb D;
Score 13.2; DB 1;
Pred. No. 2.5e+02;
4; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 13.2; DB 1;
Pred. No. 2.5e+02;
2; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 0 A; 11 C; 3 G; 4 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pavco P,
                                                             1239 GCTGGACGTGGCCATGTG 1256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Page 91; 218pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      142 CCGCTCGGCTCCGCTCCG 159
                                                                                         1 GCUGGCCGUCGCCCUGUG 18
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 1.0%;
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95US-0005974.
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                                                                                                                                                                   AAX70233 standard; RNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Escobedo J, McSwiggen J,
                                                                                                                                                                                                                               (first entry)
                               11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 72.2
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-259017/23.
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               Best Local Similarity
Matches 11; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CHIR ) CHIRON
                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-OCT-1996;
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26-OCT-1995;
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RESULT 388

The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (IL-1), kinase insert domain containing receptor (KRN) and/or foetal liver kinase 1 (flk-1) (e.g. tumour neceptor (KRN) and/or foetal liver kinase and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.

Sequence 18 BP; 0 A; 7 C; 7 G; 4 U; 0 other;

10-NOV-1997

AAT84310;

AAT84310

us09904568-3.rng

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The sequences given in AAT80211-382 represent synthetic oligonucleotides which are complementary to a portion of the 5' untranslated region (UTR) of hepatitis C virus (ECV). These sequences may be used in a pharmaceutical composition for the control or prevention of HCV infection. They may be used to inhibit replication or expression of HCV or for detecting the presence of HCV in a sample. They may be used to inhibit therefore useful in the treatment of HCV infections such as chronic and are therefor useful in the treatment of HCV infections such as chronic and acute hepatitis and hepatocarcinoma. This oligo was used in a luciferase assay to determine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligo:nucleotide(s) complementary to HCV 5' untranslated region - used in the treatment and detection of HCV infection, esp. hepatitis
                                                                Complementary; 5' untranslated region; UTR; hepatitis C virus; HCV; inhibition; replication; expression; detection; chronic hepatitis; acute hepatitis; hepatocarcinoma; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-metallothionein phosphorothioate oligodeoxyribonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.2; DB 1; Length 1
Pred. No. 2.5e+02;
0: Mismatches 3; Indels
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/note= "Phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 2 A; 3 C; 10 G; 1 T; 2 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kilkuskie RE;
Wolfe JL;
                                   Oligo HCV91, targetted to HCV region -1 to -6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         whether it binds sucessfully to its target.
                                                                                                                                                                                             7..18
/*tag= a
/note= "2'OMe modified"
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Walther DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ67556 standard; cDNA to mRNA; 18
                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                AG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HOFF ) HOFFMANN LA ROCHE & CO (HYBR-) HYBRIDON INC.
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(first entry)
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Roberts PC,
                                                                                                                                                                                                                                                1.6
/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and hepato-carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                            Key
modified_base
                                                                                                                                                                                                                                                modified base
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Roberts NA,
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15-OCT-1997
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                                                                                                                                         Synthetic.
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 ZXBXBXBXBXB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This DNA sequence comprises the junction region between intron 4 (over 10 kb) and exon 5 of the human VEGF-C gene (see also AAT84276). Exon-intron junctions were determined (see AAT84303-14) for the entire VEGF-C gene, which comprises 7 exons and 6 exons. The VEGF-C gene on chromosome 4q23 codes for a novel ligand (AAW00932) of Flt4 receptor tyrosine kinase that can be used in a claimed method to modulate growth of endothelial cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                        vascular endothelial growth factor receptor-3; ligand, angiogenesis; wound healing; lymph vessel; lymphangioma; cancer; metastasis; therapy; diagnosis; ss.
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                                                                                                                                                                                                                                                                                                                                                                  b
"5' end of exon 5, encodes amino acids
231-233 of VEGF-C"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 2.5e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                       VEGF-C; Flt4; receptor tyrosine kinase; VEGFR-3; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 4 A; 4 C; 4 G; 6 T; 0 other;
                                                                                                                    Human VEGF-C gene intron 4-exon 5 junction.
                                                                                                                                                                                                                                                                                                                                      of intron 4"
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                                                                                                                                                                                                                                                                                Location/Qualifiers
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                    ВР
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95US-0510133.
96US-0585895.
96US-0601132.
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Conservative
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                  AAT84310 standard; DNA; 18
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/*tag=
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12..18
/*tag=
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                                                                                     (first
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tes 15; Conserv
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01-AUG-1995;
12-JAN-1996;
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                                                                                                                                                                                                                                               Homo sapiens
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14-FEB-1996;

Alitalo K,

Query Match Best Local S

Matches

AAT80260

RESULT 389

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AAT80260/

13-FEB-1997

intron

exon

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Gaps

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Length 18;

PS-ODN; phosphorothioate oligodeoxyribonucleotide; chelate; ss;

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WPI; 1997-234701/21.
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                                                                                                                                                                                                               The invention relates to a new method of treating an animal suffering from heavy metal poisoning, comprising administering a phosphorothioate oligonucleotide to the animal in an amount sufficient to chelate heavy metals to cause their excretion in the urine of the animal.

The method can be used to treat poisoning caused by cadmium, lead and/or mercury. The present sequence is one of two complementary phosphorothioate oligonucleotides (PS-ODNs) which were synthesised and tested for use in the method. Each PS-ODNs wise 18 bases in length and is associated with bases 7-24 downstream from the ATG translational start site of human metallothionein II mRNA. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                   Treatment for heavy metal poisoning - by chelating heavy metal ions with phosphoro:thioate oligo:nucleotide to cause excretion in urine
                                                                                                                                                                                                                                                                                                             the anti-MT antisense sequence, while the complementary sequence is
heavy metal poisoning; cadmium; mercury; lead; metallothionein.
                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N; phosphorothioate oligodeoxyribonucleotide; chelate; ss; metal poisoning; cadmium; mercury; lead; metallothionein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Anti-metallothionein phosphorothioate oligodeoxyribonucleotide.
                                                                                                                                                                                                                                                                                                                                                              Length 18;
                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                            Score 13.2; DB 1;
Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 3 A; 3 C; 10 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ67557 standard; cDNA to mRNA; 18 BP.
                                                                                                                                                                                              Example 3; Column 10; 10pp; English
                                                                                                                                                                                                                                                                                                                                                                                                525 GCCGGAGGAGCAGCTGGG 542
                                                                                                                                                                                                                                                                                                                                                                                                                  18
                                                                                                                                                                                                                                                                                                                                                             1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                             1 GGCGCAGGAGCAGTTGGG
                                                                        91US-0759841
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91US-0759841
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              91US-0759841
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                             15; Conservative
                                                                                                              NEBRASKA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYNE-) UNIV NEBRASKA
                                                                                                                                                 WPI; 1997-234701/21.
                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                       shown in AAQ67557
                                                                                                             (UYNE-) UNIV
                                                                        12-SEP-1991;
                                                                                          12-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-AUG-1997
                                   US5618796-A
                                                     08-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US5618796-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-APR-1997
                                                                                                                                Iversen PL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Iversen PL;
                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ67557;
                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  heavy
                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 391
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ67557,
à
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The present sequence represents an oligonucleotide of the specification. The specification describes a peptide which has affinity for thrombopoietic receptor. The peptide, which can be cyclic, can be used in the preparation of therapeutics useful for treating and preventing diseases caused by abnormality in haematogenic process of megakaryocytes.
                                                                                                                                                  The invention relates to a new method of treating an animal suffering from heavy metal poisoning, comprising administering a phosphorothioate oligonucleotide to the animal in a manount sufficient to chelate heavy metals to cause their excretion in the urine of the animal.

The method can be used to treat poisoning caused by cadmium, lead and/or mercury. The present sequence is one of two complementary phosphorothicate oligonucleotides (PS-ODNs) which were synthesised and tested for use in the method. Each PS-ODNs is 18 bases in length and is associated with bases 7-24 downstream from the ATG translational sequence is shown in AAQ67556, and the present sequence is its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New recombinant therapeutic peptide - useful for, e.g. treating diseases caused by abnormality in megakaryocyte haematogenic process
Treatment for heavy metal poisoning - by chelating heavy metal ions with phosphoro:thioate oligo:nucleotide to cause excretion in urine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Thrombopoietic receptor; therapeutic drug; treatment; prevention; disease; abnormality; haematogenic process; megakaryocyte; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.0%; Score 13.2; DB 1; Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 13.2; DB 1;
Pred. No. 2.5e+02;
); Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 2 A; 10 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 1 A; 10 C; 5 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide of the specification.
                                                                                               Example 3; Column 10; 10pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 6; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ·;
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dececadeaccaerrese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             525 GCCGGAGGAGCAGCTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96JP-0231807.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV16730 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 complementary sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-234763/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JP10072492-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-MAR-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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· 0 us09904568-3.rng

(first entry)

Page 196

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PCR primer; proto-oncogene; oncogene; nucleic acid synthesis; ultrasound; stress protein; repair protein; phenylketonuria; p53 tumour suppressor; phenylalanine hydroxylase; IL-2 production; cancer; AIDS; haemophilia; autoimmune disease; chronic viral infection; cystic fibrosis; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                      Increasing nucleic acid synthesis by ultrasonic treatment of cells
                                                                                         PCR primer for the pericentrin gene.
              AAX77049 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                           (IMAR-) IMARX PHARM CORP.
                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-370731/31.
                                                                10-AUG-1999
                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                17-NOV-1997;
                                                                                                                                                                                                                                   W09925385-A1
                                                                                                                                                                                                                                                                                      11-NOV-1998;
                                                                                                                                                                                                                                                              27-MAY-1999.
                                                                                                                                                                                                Synthetic
                                       AAX77049;
                                                                                                                                                                                                                                                                                                                                                                  McCreery
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a mammalian PG1 gene and protein, and a set of PG1 biallelic markers. The PG1 polynucleotide and biallelic markers are used in a hybridisation assay, a sequencing assay, or in an allele-specific amplification assay for determining the identity of a nucleotide at a PG1-related biallelic marker. The methods can be used to detect and to assess the risk of developing cancer or prostate cancer. Barly-stage diagnosis of prostate cancer relies on prostate specific antigen (PSA) dosage. However, the effectiveness of this is limited due to its inability to discriminate between malignant and non-malignant affections of the organ, A need exists for both a reliable diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              procedure which would enable early-stage diagnosis, and for preventative and curative treatments of the disease. The PGI gene can be used for detection of prostate cancer, and the risk of developing it in the future, and can also be used to determine therapies for the disease.
                                                                                                                                                                                                                                           PGI gene; biallelic marker; PCR primer; PG1-related biallelic marker; cancer; prostate cancer; diagnosis; therapy; prostate specific antigen; PSA; human; ss.
                           Gaps
                           .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Use of a prostate cancer associated gene and biallelic markers derived from it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 2.5e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                   PCR primer for PG1 biallelic markers 4-58-289 and 4-58-318,
                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cohen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 5 A; 8 C; 3 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chumakov I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 348; 385pp; English.
                                                  703
                        0;
                                                                          8
                                                                                                                                         AAZ01197 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bougueleret L,
            83.3%;
                                                                  1 TTCGGCCCCAGCGCCCC
                                                 686 TTGGGAGCCAGCGGCCC
                                                                                                                                                                                                                                                                                                                                                                                        98WO-IB02133.
                                                                                                                                                                                                                                                                                                                                                                                                                  98US-0099658
97US-0996306
                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 1.0
Best Local Similarity 93.3
Matches 15; Conservative
                         15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-405178/34.
          Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  from it
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blumenfeld M,
                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                     WO9932644-A2
                                                                                                                                                                                                                                                                                                                                                                                        22-DEC-1998;
                                                                                                                                                                                            27-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                09-SEP-1998;
22-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                             01-JUL-1999.
                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                  AAZ01197;
                                                                                                              RESULT 393
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Unger EC;

Sadewasser D,

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98WO-US23843. 97US-0971540.

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                                                           This sequence represents a PCR primer for a proto-oncogene/oncogene, and was used to test the method of the invention. The method is for increasing synthesis of nucleic acid (I) in a cell by exposing it to ultrasound, where (I) is: (a) an endogenous sequence (Ia) encoding a stress or repair protein; or (b) an introduced exogenous sequence (Ib).
                                                                                                                                                                                                                    The method is specifically used therapeutically: (1) to treat phenylketonuria (following introduction of (1b) for phenylalamine hydroxylase); (i) to increase expression of the p53 tumour suppressor; (iii) to increase expression of the p53 tumour suppressor; (iii) to increase production of 1b.2, particularly associated with natural killer cells; and (iv) for treating cancer by administering a sequence antisense to initiation factor 3 and/or tRNA synthase. More generally, (1b) may include one or more genes or fragments, or even complete chromosomes, for delivery (in vivo, in vitro or ex vivo) to animal or plant cells for treating a very wide range of conditions, e.g. acquired immune deficiency syndrome, autoimmune diseases, chronic viral infections, haemophilia, cystic fibrosis, and cancer. Ultrasonic treatment increases expression of (I) and increases uptake of (Ib),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.0%; Score 13.2; DB 1;
33.3%; Pred. No. 2.5e+02;
Ive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 5 A; 4 C; 7 G; 2 T; 0 other;
Example 1; Page 107; 124pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        233 CTCAGGCATCTGCATCTG 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      crcaercarcreceere
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX57940;
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Gaps

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886

869 ICCCCACAGCCAAGIICC 1 rcccacacactaagagcc

18

RESULT 394 AAX77049/c

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us09904568-3.rng

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28~0CT-1999
                                                                                                                                                                                                                                                                   18
                                                                                                                                                                                                                                                                                                                            AAZ71683;
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                                                                                                                                                                                                                                  Matches
  δ
                                                                                                                                                                                                                                                                 g
                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                               This sequence represents a PCR primer for DNA encoding the D-sorbitol dehydrogenase of the invention. Cells transformed with a vector containing DNA encoding the dehydrogenase can be used to produce 1-sorbose or 2-keto-1-quionic acid as precursor for simple large-scale
               D-sorbitol dehydrogenase; L-sorbose; 2-keto-L-gulonic acid; precursor; L-ascorbic acid production; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                   Gaps
for G. oxydans D-sorbitol dehydrogenase coding sequence.
                                                                                                                                                                              Gene group for D-sorbitol dehydrogenase, useful for simple
large-scale production of L-sorbose or 2-keto-L-gulonic acid as
precursor for L-ascorbic acid
                                                                                                                                                                                                                                                                                                                   ·,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Comisky J, Fernandez J, Hoeffler J, Marcil R, Sekiguchi J;
Shuman S;
                                                                                                                                                                                                                                                                                                1.0%; Score 13.2; DB 1; Length 18; 33.3%; Pred. No. 2.5e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer PCR53 used for amplification of isolated RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Topoisomerase; 5' tagging; RNA transcript; isolating;
gene sequence; PCR primer; ss.
                                                                                                                                                 Yoshikawa
                                                                                                                                                                                                                                                                                 Sequence 18 BP; 2 A; 8 C; 3 G; 5 T; 0 other;
                                                                                                                                                Soeda S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (INVI-) INVITROGEN CORP. (SLOK ) SLOAN KETTERING INST CANCER RES.
                                                                                                                                                                                                               Example 5; Page 26; 83pp; Japanese.
                                                                                                                                                                                                                                                                                                                                 474 GGGGGAGGACTGCCGAGA 491
                                                                                                                                                Saito Y,
                                                                                                                               (FUJI ) FUJISAWA PHARM CO LID
                                                                                                                                                                                                                                                                                                                                                 degradadanaccada
                                                                                                                                                                                                                                                                                                                                                                                           ВР
                                                                                                98WO-JP04612.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98WO-US12372.
                                                                                                               97JP-0285280
                                                                                                                                                                                                                                                                                                          83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97US-0049405,
                                                                                                                                                                                                                                                                 L-ascorbic acid production.
                                                                                                                                                                                                                                                                                                                                                                                          AAX03321 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                  Conservative
                                                Gluconobacter oxydans
                                                                                                                                              Ishii Y, Noguchi Y,
                                                                                                                                                                WPI; 1999-302741/25
                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                13-OCT-1998;
                                                                                                               17-OCT-1997;
                                                               W09920763-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9856943-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                          23-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-DEC-1998
PCR primer
                D-sorbitol
                                        Synthecic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                 18
                                                                                                                                                                                                                                                                                                                                                                                                         AAX03321;
                                                                                                                                                                                                                                                                                                                                                                                RESULT 396
g
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PCR primers AAX03120-21 were used to amplify reverse-transcribed RNA alsolated using the method of the invention. The specification describes a method for covalently joining a DNA strand to an RNA strand. The method comprises forming a topolsomerase-DNA intermediate by incubating a DNA cleavage substrate comprising a topolsomerase cleavage site with a topolsomerase specific for that site, where the topolsomerase-DNA intermediate has one or more 5' single-stranded tails, and adding to topolsomerase-DNA intermediate an acceptor RNA strand complementary to DNA strand to the RNA acceptor strand and dissociation of the covalently bound topolsomerase, thereby covalently joining the DNA strand to the RNA acceptor strand and dissociation of the transcripts. They are used particularly for isolating and cloning full-length gene sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ruman biallelic marker upstream amplification primer SEQ ID NO:6039.
                          to an RNA strand, used particularly for isolating and cloning full-length gene sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human genome, biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyphing; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ..
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 13.2; DB 1;
Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 4 A; 7 C; 3 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 8; Page 1518; 2745pp; English.
                                                                                                                                                                    Disclosure; Page 63; 93pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                677 GCGTGGTATTTGGGAGCC 694
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.0%;
ilarity 83.3%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-IB00822.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GAGTCGTATATGGGAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ71683 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blumenfeld M,
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WPI; 1999-080916/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-013267/01.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    diagnosis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15;
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. 0 display peptide; screening; biomolecule activity regulator; development; liver disorder; sclerosis; cancer; hepatitis C virus; effects from pharmaceutical agents acting on a disease as well as other The present invention describes a method for screening potential regulators of the activity of a biomolecule. The method involves screening the potential regulators by selecting from a library of recombinants which express peptides at their surface, one which expresses a peptide which interacts with the biomolecule and then screening potential regulators to identify those which inhibit the interaction of the biomolecule with the selected recombinant or the peptide expressed by it. The method is used for the identification of drug molecules for treatment and prevention of diseases, especially AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: They can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side Screening biomolecule activity regulators by their effect on a biomolecule-peptide interactions for identification of potential drug Gaps treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing 0; Length 18; Score 13.2; DB 1; Length re Pred. No. 2.5e+02; Sequence 18 BP; 7 A; 0 C; 8 G; 3 T; 0 other; Oligonucleotide PCR primer SEQ ID NO:16. Example 1; Page 17; 45pp; Japanese. .0 infection; PCR primer; ss. 56 CTCCTCAATTACCCACAT 73 CTCCTCTCTATCCACAT 1 BP. 1.0%; 10-MAR-2000; 2000WO-JP01478 99JP-0063110 Eto Y; AAA92417 standard; DNA; 18 (first entry) 15; Conservative (AJIN ) AJINOMOTO CO INC. Okamoto S, Miwa K, WPI; 2000-587433/55 Query Match Best Local Similarity WO200053740-A1 10-MAR-1999; 15-JAN-2001 14-SEP-2000. Synthetic. molecules 18 Phage drug (HCV; Matches RESULT 398 AAA92417 0 D D D D D D D D D D D D D D D X X X ð g

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The present invention relates to antisense oligonucleotides (see AAAS5496-A55757) which are targeted to nucleic acids encoding a human tumour necrosis factor receptor-associated factor (TRAF). The antisense sequences comprise at least one modified internucleotide linkage, which is a phosphorothioate linkage. The oligonucleotides also include at least one modified sugar moiety such as a 2-0-methoxyethyl engar moiety. Sequences AAA55495 represent nucleotide sequences encoding human TRAF1-6. Included in the invention is a method for treating a human having a disease associated with the expression of TRAF comprising administering an antisense oligonucleotide. The reduction of jun kinase activation in cells comprises contacting the cells with an antisense oligonucleotide targeted to TRAF-2 or missense oligonucleotide targeted to TRAF-2 or TRAF-6. The antisense oligonucleotides have antiproliferative and antisense oligonucleotide targeted to TRAF-2 or TRAF-6. The antisense oligonucleotides have antiproliferative and anti-inflammatory activity and are useful for treating disorders associated with cell proliferation and inflammation. The antisense
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense oligonucleotides targeted to nucleic acids encoding human tumour necrosis factor receptor-associated factor (TRAP), useful for treating diseases associated with TRAF expression such as inflammatory diseases.
                                                                                                                                                 Gaps
of liver disorders such as sclerosis and cancer associated with hepatitis C virus (HCV) infection. The present sequence represents a PCR primer which is used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tumour necrosis factor receptor-associated factor; TRAF; human; antisense oligonuclectide; phosphorothioate; antiproliferative; anti-inflammatory; E-selectin; jun kinase; ss.
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0
                                                                                                                                                 Indels
                                                                                                            Score 13.2; DB 1;
Pred. No. 2.5e+02;
                                                                           Sequence 18 BP; 1 A; 10 C; 5 G; 2 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                            TRAF1 antisense oligonucleotide ISIS# 26699.
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0
                                                                                                                                                                                   686 TTGGGAGCCAGCGCCC 703
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                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                              1.0%;
                                                                                                                           83.3%;
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                                                                                                                                                   Conservative
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                                                                                                        Query Match
Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
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Gaps

0

Indels

Pred. No. 2.5e+02;

83.3%;

15; Conservative

Matches

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Gaps

Best Local Similarity

0; Mismatches

1300 CCTGGCCCCATGTAGCCA 1317

18

1 CCTGGTCCTGTGTAGCCA

BP.

AAA39029 standard; DNA; 18

RESULT 401 AAA39029/C AAA39029;

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The present sequence is one of a number of oligonucleotides designed to target different regions of the human NF-kappa-B p65 subunit, which is a member of the Rel/NF-kappa-B family of transcription factors.
Rel/NF-kappa-B proteins are involved in a diverse set of signaling pathways involving stress, apoptosis, cancer, growth, infection and inflammation. Antisense oligonucleotides are able to inhibit expression of the p65 subunit and may therefore be used in the treatment of disorders associated with NF-kappa-B p65 subunit expression. They may be used as a prophylaxis to prevent or delay infection, inflammation or tumour formation. Antisense compounds may also be used for research and
                                                                                                                                                                                                                                                     Human NF-kappa-B p65 subunit antisense oligodeoxynucleotide ISIS# 23738.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Autisense compounds which inhibit the expression of the human NF-kappa-B p65 subunit (p65) useful for treating diseases associated with p65 expression and as prophylaxis to prevent of delay infection,
                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= a
/note= "all or some internucleoside bonds are
/nosphozothioate and optionally some sugars may
be 2' methoxyethyl"
                                                                                                                                                                                                                                                                                 Human; anti-inflammatory; cytostatic; antimicrobial; infection;
antisense inhibition; inflammation; transcription factor;
                             ö
Length 18;
                             Indels
Score 13.2; DB 1;
Pred. No. 2.5e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 15; Column 40; 33pp; English.
                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inflammation or tumor formation
                                                           471 GCAGGGGGGGACTGCCG 488
                                                                                      1 eccedeceaedacrecre 18
                                                                                                                                                              AAA27086 standard; DNA; 18 BP.
 1.0%;
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                               15; Conservative
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                                                                                                                                                                                                                                                                                                                apoptosis; cancer; ss.
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Query Match
Best Local Similarity
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modified_base
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                                                                                                                                                                                                                          21-AUG-2000
                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US6069008-A.
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                                                                                                                                                                                             AAA27086;
                                                                                                                                  RESULT 400
                                Matches
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Bacterial, 16S rRNA; identification; polymorphism; microorganism; classification; primer; human medicine; veterinary medicine; agriculture; food science; industrial microbiology; infectious disease;

Food safety; ss

Jnidentified US6054278-A. 25-APR-2000.

Unknown bacterial 16S rRNA gene primer 0531R SEQ ID NO:7.

25-AUG-2000 (first entry)

The present invention describes a method for identifying a microorganism by comparing a composite sequence (I) of a ribosomal RNA gene region with FRAM region sequences of unknown microorganisms in a database and identifying the region in the database that matches with (I). (I) is generated by simultaneously obtaining nucleotide base sequence data from cevery copy of the KRNA gene region in the genome of the unknown microorganism. Also described is a method for identifying the species of microorganism by generating (I) and entering it into a first data register of a programmable computer, comparing the first data register with reference data registers that encode a unique composite rRNA ceptence corresponding to (I) and correlated with unique microorganism sequence corresponding first data register. The method is useful for identifying microorganisms which are useful in a variety of fields including human medicine, veterinary medicine, agriculture, food science and industrial microbiology. The microorganisms found in patients suffering from an infectious disease can also be identified.

Microorganism identification is also useful for monitoring food science also identified. The method is convenient and efficient as there is no need to isolate one or more individual 165 rRNA genes. AAA39021 to AAA39031 to examplification of the present invention. exemplification of the present invention 

Identifying an unknown microorganism by generating a composite sequence of its ribosomal RNA gene region and comparing with composite ribosomal RNA region sequences of distinct microorganisms in a database

97US-0045603.

(PEKE ) PERKIN-ELMER CORP.

Smith DH, Dodge DE; WPI; 2000-338488/29.

98US-0073465.

05-MAY-1998; 05-MAY-1997; Example; Column 10; 11pp; English

Length 18; Score 13.2; DB 1; Pred. No. 2.5e+02; Sequence 18 BP; 2 A; 7 C; 6 G; 3 T; 0 other; 1.0%; Best Local Similarity

diagnostics because they hybridise to mucleic acids encoding NP-kappa-B p65 subunit. The effect of antisense oligomucleotides on NP-kappa-B p65 subunit mRNA levels was measured using real-time quantitative PCR and Northern blot analysis. Antisense oligonucleotides were synthesised on an automated DNA synthesiser.

1.0%; Score 13.2; DB 1; Length 18;

Sequence 18 BP; 2 A; 6 C; 5 G; 5 T; 0 other;

Query Match

qq à

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to prevent infection, inflammation and tumours.
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                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                          Query Match
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                                                                                                                                                                                        RESULT 403
                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "Optionally 2'-methoxyethyl nucleotides providing bases 15..18 are also 2'-methoxyethyl nucleotides. All cytidine residues within this region are then 5-methylcytidine"
 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /noce= "Optionally 2'-methoxyethyl nucleotides
providing bases 1..4 are also 2'-methoxyethyl
nucleotides. All cytidine residues within this region are
 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense compounds targeting nucleic acids encoding human G-alpha-13 useful for treating diseases associated with G-alpha-13 expression and as prophylaxis to prevent or delay infection,
 ;
0
                                                                                                                                                                                                                                              Human, G-alpha-i3, G protein, Gi protein; adenylyl cyclase, dopamine; thyrotropin-releasing hormone; somatostatin; signal transduction pathway; antisense oligonucleotide; ss.
 Indels
                                                                                                                                                                                                                    Human G-alpha-i3 antisense oligonucleotide ISIS#25951.
                                                                                                                                                                                                                                                                                                                                                                           mod_base= OTHER
/note= "Optionally phosphorothioate
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 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                             deoxynucleotides"
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                             882 GTTCCAGGAGCTGCGGTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                         mod base=
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   Conservative
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modified_base
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   15;
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   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel antisense oligonuclectide containing compounds, useful for inhibiting the expression of G-alpha-i2 in human cells and tissues and treating infection, inflammation and cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G-alpha-i2; antisense inhibitor; infection; inflammation; prevent; tumour formation; treatment; inhibit; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              G-alpha-i2 antisense inhibitor oligonucleotide #15 (ISIS #25823).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ..
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                                                                      Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.0%; Score 13.2; DB 1; 3.3%; Pred. No. 2.5e+02; ve 0; Mismatches 3;
                                                                   1.0%; Score 13.2; DB 1;
Local Similarity 83.3%; Pred. No. 2.5e+02;
tes 15; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 4 A; 7 C; 4 G; 3 T; 0 other;
Sequence 18 BP; 5 A; 2 C; 8 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Column 40; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          877
                                                                                                                                                                                                     973 CTCACTIGACCAGICCCA 990
                                                                                                                                                                                                                                                                         Н
                                                                                                                                                                                                                                                                                                                                                                                                                                               BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GCTTTGAGGTCCCCACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GCTTTGAGGGCGTCACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0339993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0339993.
                                                                                                                                                                                                                                                                         18 crcacrrcaccrgracca
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA09715 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 1.0
Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PHARM INC
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Human; Ship-2; antisense oligonuclectide; phosphorothioate; detection;
                                                                                      inhibition; SH2-containing phosphatidylinositol phosphatase-2; ss.
                                                            Human Ship-2 phosphorothioate antisense oligonucleotide #30722.
                                                                                                                                                                                                                                                                                            Antisense oligonucleotides, useful for inhibiting human Ship-2 expression and for detecting nucleic acids encoding Ship-2 -
                                                                                                                                        /*tag= a
/note= "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                      Example 15; Column 39; 34pp; English.
                                                                                                                        Location/Qualifiers
          BP.
                                                                                                                                                                                                                      99US-0339964.
                                                                                                                                                                                                      99US-0339964.
        AAZ91440 standard; DNA; 18
                                           (first entry)
                                                                                                                                                                                                                                                        Bennett CF, Cowsert LM;
                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                      the present invention.
                                                                                                                                                                                                                                                                         WPI; 2000-181819/16.
                                                                                                                        Key
modified_base
                                                                                                       Homo sapiens
                                                                                                                                                                                                     25-JUN-1999;
                                                                                                                                                                                                                      25-JUN-1999;
                                           22-MAY-2000
                                                                                                                                                                  US6025198-A.
                                                                                                                                                                                   15-FEB-2000
                         AAZ91440;
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The present invention describes phosphorothioate antisense oligonucleotides that specifically hybridise with, and inhibit the expression of, nucleic acids encoding human Ship-2 (also called SH2-containing phosphatidylinositol phosphatase-2). Also described is a method of inhibiting the expression of Ship-2 in human cells or tissues in vitro comprising contacting the cells with the phosphorothicate antisense oligonucleotides. The phosphorothicate antisense oligonucleotides can be used to treat animals (especially humans) suspected of having or being prone to a disease or condition associated with Ship-2 expression. The present sequence represents a phosphorothicate antisense oligonucleotide for human Ship-2, from

Gaps ó Score 13.2; DB 1; Length 18; Pred. No. 2.5e+02; 3; Indels Sequence 18 BP; 2 A; 11 C; 1 G; 4 T; 0 other; 0; Mismatches 1.0%; 15; Conservative Best Local Similarity Query Match

886 <del>1</del> 869 TCCCCACAGCCAAGTICC 1 receedended

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AAZ65527 standard; DNA; 18 AAZ65527; RESULT 405 AAZ65527 ID AAZ6 XX AC AAZ6 XX XX XX XX DE Immu XX XX

BP.

Immunosuppressant inhibitor oligonucleotide TGF-betal-98-15. (first entry) 30-MAR-2000

Immunosuppressant inhibitor; transforming growth factor beta; TGF beta;

Adipocyte, hedgehog polypeptide, desert hedgehog; indian hedgehog, Ihh; sonic hedgehog; Shh; therapeutic; cytostatic; primer; RT-PCR;

PPAR-gamma mRNA amplifying RT-PCR primer R.

(first entry)

07-JAN-2002

AA166785;

ВР.

AA166785 standard; DNA; 18

RESULT 406

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This sequence is an immunosuppressant inhibitor oligonucleotide, which is used in the invention. The invention relates to a composition which contains at least one inhibitor (less than 100 kD) of a substance (e.g. transforming growth factor TGF-beta, vascular endothelial growth factor VEGF, interleukin-10 IL-10, prostaglandin E2 PGE2, or their receptors) that adversely affects the immune response. The composition also includes at least one stimulant that positively affects the immune response. This oligonucleotide is an example of an inhibitor that is used in the composition. The composition is used as an immunostimulant for the composition. The composition is used as an immunostimulant for the customent of neoplasms and infections, particularly hyperprofileration, leukaemia, (non-)Hodgkin's lymphoma; carcinoma (of oesophagus, bronchi, colon-rectum, stomach, intestine, gall bladder or duct, pancreas, anus, breast, ovary, cervix, endometrium, prostate or bladder), liver tumours, mast, of which are directed against TGFbeta or VEGF, are inhibitors of monocyte chemotactic protein-1 (MCP-1) and are useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  anti-inflammatories for treating e.g. asthma, Crohn's disease, ulcerative colitis, diabetes, glomerulonephritis, acute respiratory distress syndrome and the formation of atherosclerotic plaque.
vascular endothelial growth factor, VBGF, interleukin-10; IL-10; cancer; prosteglandin E2; PGB2; immune response; tumour; asthma; Crohn's disease; monocyte chemotactic protein-1; MCP-1; ulcerative colitis; diabetes; glomerulonephritis; acute respiratory distress syndrome; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Composition containing immune stimulant and inhibitor of agent adversely affects the immune response, for treating cancers and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%; Score 13.2; DB 1; Length 18; 33.3%; Pred. No. 2.5e+02; ve 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                            Brysch W;
                                                                                                                                                                                                                                                                                                                                                  (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 1 A; 9 C; 7 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                         Schlingensiepen R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 10; Figure 1; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               191 CCGCCCACCCGGACGCCG 208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ilarity 83.3%;
Conservative
                                                                                                                                                                                                                                             99WO-EP04013.
                                                                                                                                                                                                                                                                                      98EP-0110709.
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                                                                                                                                                                                                                                                                                                                                                                                            Schlingensiepen K,
                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-097470/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
nes 15; Conserv
                                                                                  atherosclerosis.
                                                                                                                      Unidentified
                                                                                                                                                               WO9963975-A2
                                                                                                                                                                                                                                               10-JUN-1999;
                                                                                                                                                                                                                                                                                      10-JUN-1998;
                                                                                                                                                                                                        16-DEC-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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Matches
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(CURA-) CURAGEN CORP
                                      WPI; 2001-316248/33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH75784;
                                                                                                                                                                                                                                                                                                                                                                Н
                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 408
                                                                                                                                                                                                                                                                                                                                                                                                                     AAH75784
                                                                                                                                                                                                                                                                                                                                                                                                                                            ò
                                                                                                                                                                                                                                                                                                                                                                셤
                                                                                                                                                                                                                                                                              The invention provides a method for regulating formation and/or maintenance of adipocyte tissue that comprises contacting pre adipocyte or adipocyte cells with a hedgehog polypeptide or ptc therapeutic. The method is used for regulating the growth state of an adipocyte stem/progenitor cell, and treating or preventing disorders of, or surgical or commetter repair of, adipocyte tissues, e.g. for treating or preventing hyperplastic or neoplastic conditions affecting adipocyte issue, such as soft tissue tumors, especially adipose cell tumors, e.g. lipomas, fibrollipomas, lipoblastomas, lipomatosis, hibernomas, hemangiomas and/or therapeutic agents. Sequences Ali66784-793 represent primers used in quantitative RT-PCR of PPARgamma, aP2, gli, ptc and actin mRNAs, during
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                      Method for regulating formation and/or maintenance of adipocyte tissue by contacting pre-adipocyte or adipocyte cells with a hedgehog polypeptide or ptc therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cluster, hierarchical clustering algorithm; population based study; clinical trial; DNA fingerprint; genetic profile analysis; PCR primer; SNP; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sample member clustering methed related human DNA PCR primer #20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 2.5e+02; Live 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 4 A; 5 C; 6 G; 3 T; 0 other;
                                                                                                                                                       Burtscher H;
                                                                                                                                                                                                                                                           Example; Page 76; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Greeaectrecarcreca 1
                                                                                                                                                                                                                                                                                                                                                                                                                                            the courrse of the invention.
                                                                                                                                                     Leser-Reiff U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              304 GIGGGGCIGCAACICCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-OCT-2000; 2000WO-IB01632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-OCT-1999; 99US-0161231.
07-JUL-2000; 2000US-0216897.
                                                                          28-FEB-2001; 2001WO-US06450
                                                                                                    29-FEB-2000; 2000US-186058P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF89283 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                               WPI; 2001-607352/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
les 15; Conserv
                                                                                                                           (CURI-) CURIS INC
                       WO200164238-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200129257-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GEST ) GENSET
                                                                                                                                                     Zehentner B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-APR-2001.
                                               07-SEP-2001
Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF89283;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18
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The present invention describes methods of clustering members of a sample, involving applying a hierarchical clustering algorithm to the sample members, determining the optimal number of clusters based on this and distributing the sample members into clusters using non-hierarchical clustering. The methods are useful in population based studies such as clinical trials, DNA fingerprinting appendic profile analyses. The present sequence was used to demonstrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NOV, olfactory; cytostatic; immunomodulator; vulnerary; anti-HIV; antiantalmatic; antiinflammatory; gastrointestinal; neuroprotective; osteopathic; gane therapy; odorant receptor; olfactory receptor; G-protein coupled receptor; GPCR; neuro-olfactory; trauma; PCR primer; neoplastic disorder; cancer; adenocarcinoma; lymphoma; prostate cancer; uterus cancer; immune response; AIDS; asthma; Crohn's disease; multiple sclerosis; Albright hereditary osteodystrophy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                               Genetic clustering by distributing members into optimal numbers calusters determined by a hierarchical clustering algorithm or by paired-pair analysis of homozygous pairs in clusters got from non-hierarchical clustering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%; Score 13.2; DB 1;
33.3%; Pred. No. 2.5e+02;
ive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 5 A; 8 C; 3 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                          Claim 61; Page 78; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human NOV 12 reverse PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18
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27-JAN-2000; 2000US-0178414.
07-FEB-2000; 2000US-0180634.
24-JUL-2000; 2000US-0220516.
28-JUL-2000; 2000US-0221408.
31-JUL-2000; 2000US-0221408.
21-DEC-2000; 2000US-02599.
08-JAN-2001; 2001US-0260290.
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27-JAN-2000; 2000US-0178406.
27-JAN-2000; 2000US-0178408.
27-JAN-2000; 2000US-0178409.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         869 TCCCCACAGCCAAGTTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TCCCCACAGCTAAGAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JAN-2001; 2001WO-US02849
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAH75784 standard; DNA; 18
m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-OCT-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match
Best Local Similarity 83.3
Matches 15; Conservative
Schork N, Skierczynski
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The present invention relates to novel human NOVX proteins and coding agequences, where X is any number from 1 to 18 (see AAH75716-AAH75733, and AAG64400 and AAG66322-AAG66338). NOVX are members of the charactery receptor (OR) family, which are G-protein coupled receptors (GPCRs). The NOVX proteins and coding sequences are useful as therapeutics, particularly in the manufacture of a medicament for treating a syndrome associated with a human diseases/disorders of the neuro-offactory system, e.g. those induced by trauma, surgery and/or neoplastic disorders. Furthermore, the coding sequences and proteins are useful in treating cancer e.g. adenocarcinoma, lymphoma, prostate cancer, ulterus cancer, inappropriate immune response, AIDS, asthma, Crohn's disease, multiple sclerosis or Albright hereditary ottodystrophy. The coding sequences are also useful in hereditary ottodystrophy. The coding sequences are also useful in easing the above present income. The present PCR primer was used in an example from the
Spytek KA, Li L, Tchernev VT, Vernet CAM;
                                                                                                                             New NOVX polypeptides and polynucleotides, useful for treating or preventing a syndrome associated with a human disease (e.g. disorders of the neuro-olfactory system), as well as in gene therapy -
                                                                                                                                                                                                                                       Example 2; Page 229; 242pp; English.
  Padigaru M,
Macdougall J;
                                                                              WPI; 2001-514556/56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present invention.
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Sequence 18 BP; 5 A; 5 C; 7 G; 1 T; 0 other;

; 0 1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 2.5e+02; 3; Indels 0; Mismatches 83.3%; Local Similarity 83.3 es 15; Conservative Query Match Matches

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AAH51027 standard; DNA; 18 BP 28-AUG-2001 AAH51027; RESULT 409 AAH5102 

(first entry)

Human nGPCR9 PCR primer #2.

G protein-coupled receptor; nGPCR; seven transmembrane receptor; signal transduction; schizophrenia; thyroid disorder; renal failure; rheumatoid arthritis; CNS disorder; infection, metabolic disease; cardiovascular disease; proliferative disorder; hormonal disorder; neurological disorder; neuronal disorder; Alzheimer's disease; attention deficit-hyperactivity disorder/attention deficit disorder; Parkinson's disease; migraine; senile dementia; inflammatory disease; rheumatoid arthritis; autoimmune disorder; respiratory ailment; neuroprotective; PCR primer; ss.

Homo sapiens

WO200136473-A2.

25-MAY-2001

16-NOV-2000; 2000WO-US31581.

99US-0166071. 99US-0166678. 99US-0173396. 2000US-0184129. 2000US-0185421. 2000US-0185554. 22-FEB-2000; 28-FEB-2000; 28-FEB-2000; 19-NOV-1999; 28-DEC-1999; 16-NOV-1999; 17-NOV-1999;

2000US-0190310. 2000US-0190800. 2000US-0198568. 2000US-0201190. 2000US-0203111. 2000US-0186811. 2000US-0188114. 25-MAY-2000; 2000US-0207094 20-APR-2000; 2 02-MAY-2000; 2 08-MAY-2000; 2 21-MAR-2000; 03-MAR-2000; 17-MAR-2000; 

(PHAA ) PHARMACIA & UPJOHN CO

Wood LS, Parodi LA, Hiebsch RR, Lind P, Slightom J; A, Kaytes PS, Bannigan CM, Ruff V, Sejlitz T, Huff RM; Schellin KA, Vogeli

WPI; 2001-389826/41.

New G protein-coupled receptor (nGPCR-x) and its encoding polynucleotide useful for diagnosing and treating e.g. schizophrenia

Example 4; Page 116; 261pp; English.

Sequence 18 BP; 1 A; 5 C; 8 G; 4 T; 0 other;

; 0 Score 13.2; DB 1; Length 18; Pred. No. 2.5e+02; Indels 0; Mismatches 1.08; 83.3%; Best Local Similarity 83.3 Matches 15; Conservative Query Match

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RESULT 410 AAF26667

AAF26667 standard; DNA; 18 BP.

(first entry) 02-APR-2001

Human Smad7 phosphorothioate antisense oligonucleotide SEQ ID NO:10. 

Human, Smad7; antisense oligonucleotide; phosphorothioate; inhibition; antiinflammatory; cytostatic; infection; inflammation; tumour formation;

sapiens. Ношо

Key

Location/Qualifiers

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Antisense compound capable of inhibiting the expression of human Smad7, useful for preventing or delaying infection, inflammation or tumor formation -
                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes an antisense compound (I) of up to 30 nucleobases in length capable of inhibiting the expression of human Smad7. (I) has antiinflammatory and cytostatic, and is a modulator of smad7 expression. (I) can be useful for inhibiting the expression of human Smad7 in human cells or tissues, in vitro. (I) is commonly used as a research reagent and in diamostics for example, to elucidate the function of particular genes. (I) is also useful for distinguishing between functions of various members of a biological pathway and for research use. (I) is also utilised for diagnostics, therapeutics, prophylaxis and in kits. (I) is also useful prophylactically, e.g. to prophylaxis and in kits. (I) is also useful prophylactically, e.g. to prevent or delay infection, inflammation or tumour formation. AMP26667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Vascular endothelial growth factor receptor 3 inhibitor; VEGFR-3; angiogenesis; lymphangiogenesis; vascular endothelial growth factor; cytostatic; hepatotroppt; antiinflammatory; hypotensive; antidiabetic; vulnerary; cell surface receptor; cancer; neovascularisation; liver disease; hypertension; post-trauma; chronic hepatitis; primer; haemangioma; diabetes; PDGF; platelet derived growth factor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            to AAF26706 represent human Smad7 antisense oligonucleotides from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VEGFR-3 binding peptide library related primer SEQ ID NO:75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 13.2; DB 1; Length 18; Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3; Indels
               /*tag= a
/note= "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 1 A; 12 C; 3 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Column 40; 33pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              583 CTCCGTCTGCCCCCACC 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.0%;
                                                                                                                               09-JAN-2000; 2000US-0487444
                                                                                                                                                                 09-JAN-2000; 2000US-0487444
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 83.3
                                                                                                                                                                                                   PHARM INC
                                                                                                                                                                                                                                  Cowsert LM;
                                                                                                                                                                                                                                                                  WPI; 2001-070108/08.
modified base
                                                                                                                                                                                                 SISI (-SISI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-JAN-2003
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                                                               US6159697-A.
                                                                                               12-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                  Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABQ82729;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 411
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The present invention describes an isolated peptide (1) that binds to and inhibits vascular endothelial growth factor receptor 3 (VBGFR-3).

(1) have cytostatic, hepatotropic, antiinflammatory, hypotensive, antidiabetic and vulnerary activities, and can be used in gene therapy. Compositions and methods from the present invention are useful for diagnosing, evaluating and treating disorders mediated by the activity of the cell surface receptor VBGFR-3 such as cancer, e.g. brain, lung, liver, spleen, kidney, lymph node, small intestine, blood cells, pancreas, colon, stomach, breast, endometrium, prostate, resticle, ovary, skin, head and neck, oeosphagus, bone, marrow or blood, and diseases of neovascularisation, e.g. liver diseases, hypertension, copst-trauma, chronic hepatitis, haemangiomas and diabetee. The present sequence represents a primer used in the construction of a VBGFR-3 binding peptide library, which is used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nG protein coupled receptor; communication; serpentine structure; seven transmembrane receptor; 7TM; mental disorder; diagnosis; genetic predisposition; brain; immune response; gene therapy; anxiety disorder; depression; bipolar disorder; schizophrenia; Huntington's disease; dyskinesia; manic depression; stroke; Parkinson's disease; Alzheimer's disease; disbates; inflammation; wound;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                          useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ss; G protein-coupled receptor; GPCR; nGPCR; beGPCR;
                                                                                                                                                                                                         diagnosing, evaluating, treating disorders mediated by VEGFR-3 activity, such as cancer and diseases of neovascularization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3; Indels
                                                                                                                                                                                          New isolated peptide that inhibits VEGF-C and VEGF-D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer, #6, used to detect expression of nGPCR-9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%; Score 13.2; DB 1;
83.3%; Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 1 A; 10 C; 5 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                           Example; Page 74; 149pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     686 TTGGGAGCCAGCGCCC 703
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP.
16-JAN-2002; 2002WO-IB00099.
                                17-JAN-2001; 2001US-262476P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-FEB-2001; 2001WO-US04641.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 83.3°
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-NOV-2002 (first entry)
                                                                                                                       Koivunen E,
                                                                                    (LICH ) LICENTIA LTD.
                                                                                                                                                        WPI; 2002-691521/74.
                                                                 (LUDW-) LUDWIG INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200264789-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-AUG-2002
                                                                                                                     Alitalo K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention
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WO200257299-A2

25-JUL-2002

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Query Match
Best Local Similarity
                                                                                                                            US2002081577-A1.
                                                                                                                                    02-JUL-1997;
                                                                                                        15-NOV-2002
                                                                                                                                27-JUN-2002
                                                                                                                         Synthetic
                                                                                                    ABS65844;
   Lind P,
                                                                                              RESULT 413
                                                                                Matches
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97US-0887505.

11-DEC-2000; 2000US-254669P. 19-SEP-2001; 2001US-323400P. WOLFE J L. ROBERTS P C. HAMLIN H A. FRANK B L. GOODCHILD J. ROBERTS N A. WALTHER D M. WPI; 2002-537132/57. WO200248401-A2. 28-OCT-2002 02-JUL-1996; 06-JUN-1995; Homo sapiens 20-JUN-2002 18 ABT06049; Query Match (KILK/) (FRAN/) (GOOD/) (WALT/) (HAML/) (ROBE/) [ROBE/] RESULT 414 ABT06049, ⋧ 8 ÷ New nucleic acids and polypeptides of the nG protein-coupled receptor, useful for treating or diagnosing a mental disorder or a disorder affecting the brain, e.g. anxiety disorders, schizophrenia, stroke or Parkinson's disease primers which were used to amplify, and detect, the DNA encoding the nGPCRs (also referred to as beGPCRs). Gaps Hepatitis C virus; HCV; hepatocyte infection; non-A hepatitis; non-B hepatitis; acute hepatitis; chronic hepatitis; hepatocellular carcinoma; virucide; cytostatic; antisense therapy; gene therapy; ss; DNA-RNA hybrid. ; Inhibitory oligonucleotide specific for hepatitis C virus #50. 1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 2.5e+02; ive 0; Mismatches 3; Indel8 Sequence 18 BP; 1 A; 5 C; 8 G; 4 T; 0 other; Wood LS Example 4; Page 110; 244pp; English. 524 TGCCGGAGGAGCAGCTGG 541 18 Vogeli G, ABS65844 standard; DNA; 18 BP. (PHAA ) PHARMACIA & UPJOHN CO TGCCTGTGGAGCCGCTGG (first entry) 15; Conservative Parodi LA, WPI; 2002-674879/72

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The invention describes synthetic oligonucleotides complementary to a portion of the 5' untranslated region of hepatitis C virus. The oligonucleotides may be used in methods for controlling, preventing, and treating hepatitis C virus infection, in antisense technology and gene therapy, and of detecting the presence of hepatitis C virus in a sample. Hepatitis C virus (HCV) is an enveloped, positive sense, single-stranded RNA virus which infects hepatocytes. HCV is the major cause of non-A, non-B, acute and chronic hepatitis, and has been associated with hepatocellular carcinoma. The invention describes methods and kits for inhibiting replication of HCV, inhibiting the expression of HCV nucleic acid and protein, and for treating HCV infections. This sequence represents a synthetic DNA-RNA hybrid oligonucleotide used for inhibiting HCV replication and expression of HCV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Single Primer Amplification; nested oligonucleotide extension reaction; hairpin; SPA; library; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic oligonucleotides complementary to a portion of the 5' untranslated region of hepatitis C virus (HCV), useful for diagnosing and treating HCV infections and hepatocellular carcinoma -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                               Roberts PC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%; Score 13.2; DB 1; Length 18; 33.3%; Pred. No. 2.5e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human IgM heavy chain gene related PCR primer SEQ ID No 63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 2 A; 3 C; 10 G; 1 T; 2 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Wolfe JL,
                                                                                                                                                                                                                                                                                                                                                                                                                               Goodchild J,
Walther DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 22; Page 11; 74pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          631 CICCAGGAGCICIGCAIC 648
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                                     95US-0471968.
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96US-021104P.
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Matches 15; Conservative
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                                                                                                        KILKUSKIE R L.
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16:51:54 2004

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Thu Jan

WPI; 2002-315546/35.

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(ALEX-) ALEXION PHARM INC
                                                                                             WO200218616-A1
                                                                                                07-MAR-2002.
                                                                            18-JUN-2002
                                                                                                         (HITB )
(HITB )
                                                                                                              Ke S;
                                                                  RESULT 415
                                                                    ABK47739,
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The invention relates to a method for amplifying a nucleic acid using Single Primer Amplification (SPA). The method comprises synthesising a Single Primer Amplification (SPA). The method comprises synthesising a structure with the nested oligomucleotide extension reaction. The method is useful for amplifying a nucleic acid, preferably for amplifying a prolypeptides encoded by the sequences to build a complex library of polypeptides encoded by the sequences. The engineered nucleic acid strand is useful for amplifying a nucleic acid strand by providing a nucleic acid with a predetermined sequence engineered nucleic acid strand sequence complementary to the predetermined sequence and a hairpin structure between them and contacting the engineered nucleic acid strand with a primer containing at least a portion of the predetermined sequence acomplementary nucleic acid strand. The method of the invention is useful complementary nucleic acid strand. The method of the invention is useful for producing large amounts of a target nucleic acid sequence and for amplifying simultaneously more than one different target nucleic acid sequence located on the same or different nucleic acid molecules. This polymucleotide sequence represents a PCR primer of the invention.
                                                                                                                                                                                         Amplifying nucleic acid by synthesizing template nucleic acid containing a predetermined sequence and hairpin structure and using the template for target amplification by Single Primer Amplification -
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Maruyama
Lin Y, Mowhirter J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 1 A; 2 C; 7 G; 8 T; 0 other;
Bowdish KS, Barbas-frederickson S,
                                                                                                                                                                                                                                                                                                                                                                                                                             54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 22;
                                                                                                       WPI; 2002-500537/53
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0 Gaps . 0 Score 13.2; DB 1; Length 18; Pred. No. 2.5e+02; 0; Mismatches 3; Indels 0 1.0%; Conservative Query Match Best Local Similarity Matches 15; Conserv

ABK47739 standard; DNA; 18 BP (first entry)

Beta-actin reverse PCR primer used in invention relating to TMOs

Primer-dependent polymerase-mediated DNA synthesis; TMO; template-mimic oligonucleotide; nucleic acid amplification; multiplex RT-PCR; reverse transcriptase-PCR; Competimer method; primer; beta-actin; ss.

Unidentified.

30-AUG-2001; 2001WO-US27287

01-SEP-2000; 2000US-230184P.

HITACHI CHEM CO LTD. HITACHI CHEM RES CENT INC.

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The present invention relates to a method of modulating amplification efficiency of a target sequence in primer-dependent polymerase-mediated DNA synthesis. The method comprises adding a template-mimic oligonuclectide (TMO) to a primer-dependent polymerase-mediated DNA synthesis reaction mixture containing primers, to block a primer for amplifying a target sequence in the mixture from hybridising to the target sequence. The method is useful for adjusting the efficiency of target template nucleic acid amplification by controlling the ratio of template-like oligonucleotides to amplification primers. The new method provides a convenient and efficient method for simultaneous detection of high and low-abundant genes in multiplex RT-PCR, and is more potent and easier to control than the Competimer method. The present sequence represents a PCR primer used in a real-time PCR assay in the example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; immunosuppressive; antirheumatic; antiarthritic; antiulcer; antianaemic; antipsoriatic; B7-related polypeptide; BSL1; BSL2; BSL3; autoimmune disease; rheumatoid arthritis; multiple sclerosis; Hashimoto's thyroiditis; Graves' disease; Crohn's disease; psoriasis; ulcerative colitis; permicious anaemia; bone marrow transplantation; graft versus host disease; organ transplantation; BSR.
                                                                                         primer-dependent polymerage-mediated DNA synthesis, useful for adjusting the efficiency nucleic acid amplification comprises adding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                         Modulating amplification efficiency of a target sequence in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 18;
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Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 2 A; 7 C; 3 G; 6 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  B7-related protein, BSL2, PCR primer #5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8
                                                                                                                                                                                                                                     Example; Page 12; 39pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   935 TGGAGAAGAGGTGTGAGC 952
                                                                                                                                                                 template mimic-oligonucleotide
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Best Local Similarity 83.3%;
Matches 15; Conservative (
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28-FEB-2001; 2001US-272107P.
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The invention relates to novel nucleic acids encoding B7-related polypeptides. The B7-related polypeptides include the BSL1, BSL2, or BSL3 polypeptides. The mucleic acid, polypeptide, and and antibodies are useful for treating autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, Hashimoto's thyroiditis, graves' disease, Crohn's disease, ulcerative collitis, pernicious anaemia and psoriasis. They may also be used to treat tissue, bone marrow, and organ transplantation, and graft versus host disease. ABK24010-ABK24093 regresent B7-related proteins, BSL1, BSL2 and BSL3 coding sequences and
Example 3; Page 101; 179pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   represent B7-related proteins, PCR primers of the invention.
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Score 13.2; DB 1; Length 18; Pred. No. 2.5e+02; 0; Mismatches 3; Indels Sequence 18 BP; 4 A; 7 C; 3 G; 4 T; 0 other; 1014 CCTGAGATGGTGCCAAAG 1031 cererearegreacagae 1 1.0%; 15; Conservative Query Match Best Local Similarity Matches

18 g Š

ABL43688 standard; DNA; 18 RESULT 417

BP

Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; Human chromosome 1p36-35 PCR primer SEQ ID NO:732 11-APR-2002 ABL43688; ABL43688

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 AGC
 ABL4

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 ABL4

XX
 ABL4

XX
 ABL6

XX
 BUMB

X

(first entry)

12-MAR-2001; 2001JP-0068285. genome; PCR primer; ss. JP2001321190-A. Homo sapiens. 20-NOV-2001.

(RIKA ) RIKAGAKU KENKYUSHO. (GENO-) GENOTEX YG.

10-MAR-2000; 2000JP-0066716.

WPI; 2002-144136/19.

Claim 4; Page 19; 528pp; Japanese. Arraying genome clones

resultant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant applates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell discrimination Nos. to mixed clones in the multiwell plates of the specified and alternal directions; (f) the mixed clones are cultured and the

reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABL42957 to ABL45222 represent PCR primers for human chromosome 1936-35 DNA, and ABL45323 to ABL45634 represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention. 88888888

Sequence 18 BP; 5 A; 7 C; 4 G; 2 T; 0 other;

ò Gaps .. 0 Length 18; Indels Score 13.2; DB 1; Pred. No. 2.5e+02; 0; Mismatches 3; Query Match Best Local Similarity 83.3%; Matches 15; Conservative

559 ATGCACACACTGCTCCAG 576 18 1 AAGGCCACACTGCTCCAG 원 ઠે

ABT21516 standard; DNA; 18 RESULT 418 ABT21516/

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Gaps

; 0

BD.

ABT21516; 

16-APR-2003 (first entry)

Racing potential; horse; grandpaternal DNA; over-represented; breeding; grandmother; performance; progeny horse; PCR; primer; ss. Multiplex group PCR primer #263.

Unidentified.

WO200292851-A2.

21-NOV-2002.

15-MAY-2002; 2002WO-GB02273.

15-MAY-2001; 2001GB-0011886.

(ANIM-) ANIMAL HEALTH TRUST. (BRHO-) BRITISH HORSERACING BOARD.

Binns MM, Swinburne JE;

WPI; 2003-129314/12.

Determining the racing potential of a horse comprises measuring whether grandpaternal or grandmaternal DNA from the selected grandmother DNA is over-represented in the genome of the horse

Example 2; Page 25; 49pp; English

The invention relates to a novel method for determining racing potential of a horse. The method comprises measuring: whether grandpaternal DNA is over-represented in the genome of the horse; or in the case where one of the grandmothers was selected for breeding on the basis of racing performance, whether grandmaternal DNA from the selected grandmother is over-represented in the genome of the horse which indicates that the horse has good racing potential. The method of the invention is useful for determining the racing potential of a horse or for obtaining a progeny horse with good racing potential of a horse or for obtaining a progeny horse with good racing potential. This polynucleotide sequence represents a PCR primer used in the detection method of over-representation of DNA from male grandparents of the invention.

Seguence 18 BP; 2 A; 12 C; 2 G; 2 T; 0 other;

Gaps . 0 1.0%; Score 13.2; DB 1; Length 18; 33.3%; Pred. No. 2.5e+02; ive 0; Mismatches 3; Indels 83.3%; 15; Conservative Local Similarity Query Match Matches

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528 GGAGGAGCAGCTGGGTGC 545

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Thu Jan

18 GGTGGAGCAGGTGGGGGC

Early mitotic inhibitor 1; Emil; anaphase-promoting complex; APC; mitosis; cell cycling; hyperproliferative condition; oocyte activation; gene therapy; PCR; primer; ss.

PCR primer for human early mitotic inhibitor 1 (Emil) cDNA.

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The invention comprises the amino acid and coding sequence of B7-related (BSL2) fusion proteins. The B7-related fusion proteins of the invention are useful for modulating the activation of immune or inflammatory response cells (e.g. T cells). The B7-related fusion proteins are useful for treating or preventing: transplantation rejection, graft versus host disease, asthma; chronic obstructive pulmonary disease, cancers; viral infections (e.g. HIV, herpes or encephalitis); and autoimmune disease (e.g. rheumatoid arthritis, multiple sclerosis or psoriasis). The present by sequence represents a PCR primer that was used in an example of the
                                                                                                                                        PCR, ss; gene therapy, B7-related fusion protein; BSL2; viral infection; immune response modulation; inflammatory response modulation; cancer; transplantation rejection; graft versus host disease; asthma; herpes; chronic obstructive pulmonary disease; HIV; encephalitis; psoriasis; autoimmune disease; rheumatoid arthritis; multiple sclerosis; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated B7-related nucleic acid fusion molecules and fusion polypeptides, useful for diagnostic applications, modulating the activation of immune or inflammatory response cells, preventing of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13.2; DB 1; Length 18;
Pred. No. 2.5e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 4 A; 7 C; 3 G; 4 T; 0 other;
                                                                                                           B7-related PCR primer - SEQ ID No 21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 129; 188pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BRIM ) BRISTOL-MYERS SQUIBB CO.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treating cancer or psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                           06-JUN-2001; 2001US-0875338.
15-FEB-2002; 2002US-0077023.
                                                                                                                                                                                                                                                                                                                                                                        06-JUN-2002; 2002WO-US18049.
ABT15904 standard; DNA; 18
                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shen H;
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Best Local Similarity
                                                                                                                                                                                                                                                                                              WO200299119-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mikesell GE,
                                                                         28-MAR-2003
                                                                                                                                                                                                                                                                                                                                   12-DEC-2002
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                                   ABT15904;
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pCR primers ABV77248-49 were used to amplify CDNA encoding human early mitotic inhibitor 1 (Emil) polypeptide. Emil polypeptides are used in the method of the invention to inhibit the anaphase-promoting complex (APC) in a proliferating cell. To this end, the Emil polypeptide is provided to a cell undergoing mitosis. The method is used to inhibit APC in a proliferating cell. Emil is used to screen for biologically active agents that modulate Emil function or that mimic Emil activity, and the agents are used for enhancing APC in a proliferating cell. The Emil is useful for modulating the cycling of cells, e.g. for treating hyperproliferative conditions, in diseases involving tissues where there is a high rate of cell turnover, and in modulating ocyte activation.

Inhibiting the anaphase-promoting complex in a proliferating cell, useful for treating hyperproliferative conditions, comprises providing an early mitotic inhibitor 1 polypeptide to a cell undergoing mitosis

Example 2; Page 42; 82pp; English.

(STRD ) UNIV LELAND STANFORD JUNIOR.

Reimann JDR;

Jackson PK,

WPI; 2003-129363/12.

23-MAY-2002; 2002WO-US16346. 24-MAY-2001; 2001US-293921P.

WO200294198-A2.

28-NOV-2002

Homo sapiens

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determining specific analogues, agonists, antagonists and mimetics. Nucleic acid compositions are useful in identifying homologous or related genes, in producing the encoded protein, in producing compositions that modulate the expression or function of its encoded protein, for gene therapy, mapping functional regions of the protein, and in studying
                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DPP10; dipeptidyl peptidase; prolyloligopeptidase; enzyme; asthma; antiinflammatory; antiasthmatic; antipsoriatic; antiarthritic; antirheumatic; vaccine; gene therapy; inflammatory disease; antirheumatory bowel disease; atopy, rheumatord arthritis; psoriasis;
                                                                                                                                                                                                                                                                                                                         .
0
                                                                                                                                                                                                                                                                  Length 18;
                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                            Score 13.2; DB 1;
Pred. No. 2.5e+02;
0; Mismatches 3;
                                                                                                                                                                                                            Sequence 18 BP; 5 A; 1 C; 10 G; 2 T; 0 other;
                                                                                                                                                        associated physiological pathways.
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                                                                                                                                                                                                                                                                     1.0%;
                                                                                                                                                                                                                                                                                              83.3%;
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                                                                                                                                                                                                                                                                                              Best Local Similarity
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                                                                                                                                                                                                                                                                     Query Match
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ABQ84276/c
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Gaps

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1014 CCTGAGATGGTGCCAAAG 1031

15; Conservative

Matches

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ccrercardercadada

18

ABV77248 standard; DNA; 18

RESULT 420

ABV77248

(first entry)

28-MAR-2003

ABV77248;

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0;

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The present invention describes a new isolated nucleic acid sequence (I) comprising a DPP10 mRNA sequence. DPP10 is a dipeptidyl peptidase (also known as prolyloligopeptidase). (I) has antiinflammatory, antiasthmatic, antiasthmatic, antiasthmatic, antistendatic, antiasthmatic, antiporatic, and antitheumatic activities, and can be used in vaccines and gene therapy. A composition comprising (I) can be used for the manufacture of a medicament for regulating DPP10 expression or for preventing or treating inflammatory disease e.g., inflammatory bowel disease, asthma, atopy, rheumatoid arthritis or psoriasis. (I) can also be used in an assay for detecting or measuring DPP10 in a sample. A host cell comprising (I) can be used for producing recombinant DPP10 gene products, or in drug screening systems to identify agents for diagnosis or treatment of individuals having or susceptible to inflammatory disease. Human DPP10 is located on chromosome 2, more specifically chromosome 2414. ABQ84524 to ABQ84612 and ABPS5569 to ABDS5629 represent sequences used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                 New nucleic acid sequence comprising DPP10 mRNA, useful for the manufacture of a medicament for regulating DPP10 protein expression or for preventing or treating inflammatory disease e.g., inflammatory bowel disease
                                                                                                                                                                                                                                                                                                                                                               Lench N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 2 A; 7 C; 3 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                               Cookson WOCM, Moffat MF, Allen M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; Page 70; 321pp; English.
  chromosome 2q14; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.0%;
                                                                                                                                                                                                                    24-APR-2001; 2001GB-0010044.
24-APR-2001; 2001GB-0010046.
12-OCT.-2001; 2001GB-0024575.
12-OCT-2001; 2001GB-0024594
                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS INNOVATIONS LTD
                                                                                                                                                                              24-APR-2002; 2002WO-GB01887.
                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-093132/08.
                                                                                                WO200286113-A2.
                                        sapiens.
                                                                                                                                        31-OCT-2002
                                                      Synthetic.
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Best Local 9
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                              Gaps
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Score 13.2; DB 1; Length 18;
Pred. No. 2.5e+02;
0; Mismatches 3; Indels
                           0
                                                      935 TGGAGAGAGGTGTGAGC 952
                                                                             18 regadadadadcraceaec 1
               l Similarity 83.3
15; Conservative
                            Matches
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Gaps

Haematopoietic cell proliferation disorder related oligonucleotide #585. ABZ10445 standard; DNA; 18 BP (first entry) 16-JAN-2003 ABZ10445; RESULT 422 ABZ10445/ ID ABZ1 XX ABZ1 XX ABZ1 XX I6-J XX I6-J XX KW Humë KW Gene KW Gytc XX Cytc XX Cytc XX OS Homc OS Synti

Human; haematopoietic cell proliferation disorder; cytostatic; gene therapy; lymphocytic leukaemia; acute myelogenous leukaemia; cytosine methylation state; probe; primer; ss.

Homo sapiens.

Synthetic,

Human; haematopoietic cell proliferation disorder; cytostatic; gene therapy; lymphocytic leukaemia; acute myelogenous leukaemia; cytosine methylation state; probe; primer; ss.

The present invention describes a method for detecting and differentiating between haematopoietic cell proliferative disorders associated with at least 1 gene and/or their regulatory regions in a subject. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least 1 reagent, which distinguishes between methylated and non-methylated CpG minucleotides within the target nucleic acid. ABZ09661 to ABZ1018 represent specifically claimed nucleotide sequences from the present invention. Oligonucleotides from the present invention on be used: for differentiating between healthy haematopoietic cells and proliferative disorder haematopoietic cells; for differentiating between acute sphotosytic leukaemia and acute myelogenous leukaemia; as probes for determining the cytosine methylation state and/or single nucleotide polymorphisms (SNPS) of haematopoietic cell proliferation disorder related only haematopoietic cell proliferation disorder related on DNA sequences and their complements; and as primers for the manaloguesic proliferation of sequences from the present invention can also be used for detecting a predisposition to, differentiation between subject haematopoietic cell proliferative disorders. The present method enables hamply specific classification of haematopoietic cell proliferative disorders allowing for improved and informed treatment of patients. Haematopoietic cell proliferation disorder related oligonucleotide #586. Detecting and differentiating between hematopoietic cell proliferative disorders, comprises contacting a target nucleic acid with a reagent that distinguishes between methylated and non-methylated CpG Berlin K, Braun A, Distler J, Guetig D, Howe A, Mueller J; Olek A, Piepenbrock C, Adorjan P, Grabs G, Lesche R, Leu E; Lewin A, Lipscher E, Maier S, Model F, Mueller V, Otto T; Pelet C, Schwope I, Ziebarth H; . 1.0%; Score 13.2; DB 1; Length 18; 33.3%; Pred. No. 2.5e+02; 3; Indels Sequence 18 BP; 1 A; 1 C; 8 G; 8 T; 0 other; Mismatches Claim 15; SEQ ID 585; 117pp; English. . 0 24 AACCAAACCCAGCTACGC 41 Н 26-MAR-2002; 2002WO-EP03401. 18 AACCAAACCGACCTACAC 26-MAR-2001; 2001US-278333P. 83.3%; ABZ10446 standard; DNA; 18 (first entry) Best Local Similarity 83.3 Matches 15; Conservative (EPIG-) EPIGENOMICS AG. WPI; 2003-018942/01 WO200277272-A2 dinucleotides 16-JAN-2003 03-OCT-2002 ABZ10446; Query Match ABZ10446/ d ð

us09904568-3.rng

cytosine methylation state; probe; primer; ss.

Thu Jan

sapiens

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ABZ11020,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a method for detecting and differentiating between haematopoietic cell proliferative disorders associated with at least 1 gene and/or their regulatory regions in a subject. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least 1 reagent, which distinguishes between methylated and non-methylated CpG dinucleotides within the target nucleic acid. ABZ09861 to ABZ1118 represent specifically claimed nucleotide sequences from the present invention. Oligonucleotides from the present invention can be used: for invention. Oligonucleotides from the present invention can be used: for differentiating between healthy haematopoietic cells and proliferative disorder haematopoietic cells; for differentiating between acute solventianing the cytosine methylation state and/or single nucleotide polymorphisms (SNPs) of haematopoietic cell proliferation disorder related sequences and their complements; and as primers for the maplification of haematopoietic cell proliferation disorder related mapping a predisposition to, differentiation between and a primers for the amplification of haematopoietic cell proliferation disorder related between the present invention can be used for detecting a predisposition to, differentiation between
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            subclasses, diagnosis, prognosis, treatment and/or monitoring of haematopodetic cell proliferative disorders. The present method enables a highly specific classification of haematopoietic cell proliferative disorders allowing for improved and informed treatment of patients.
                                                                                                                                                                                                                                                                                                                                                                                                                  Detecting and differentiating between hematopoietic cell proliferative disorders, comprises contacting a target nucleic acid with a reagent that distinguishes between methylated and non-methylated CpG dinucleotides -
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gene therapy; lymphocytic leukaemia; acute myelogenous leukaemia;
                                                                                                                                                                                                                                                                      Mueller J;
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                                                                                                                                                                                                                                                                    , Braun A, Distler J, Guetig D, Howe A, Mueller J, Piepenbrock C, Adorian P, Grabs G, Lesche R, Leu E Lipscher E, Mater S, Model F, Mueller V, Otto T; Schwope I, Ziebarth H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 2.5e+02; iive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 1 A; 0 C; 8 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 15; SEQ ID 586; 117pp; English.
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Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                               (EPIG-) EPIGENOMICS AG.
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                                                             WO200277272-A2.
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Pelet C,
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                        Synthetic.
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The present invention describes a mechoa for detecting and present invention describes a methoa for detecting and associated with at least 1 gene and/or their regulatory regions in a subject. The method comprises contacting a target mucleic acid in a biological sample obtained from the subject with at least 1 reagent, which distinguishes between methylated and non-methylated CpG which distinguishes between methylated and non-methylated CpG dinucleotides within the target nucleic acid. ABZOSSE1 to ABZOSSE1 to ABZILLIS represent specifically claimed nucleotide sequences from the present invention. Oligonucleotides from the present invention can be used: for differentiating between healthy haematopoietic cells and proliferative disorder haematopoietic cells, for differentiating between acute lymphocytic leukaemia and acute myelogenous leukaemia; as probes for determining the cytosine methylation state and/or single nucleotide collymophymorphisms (SNPs) of haematopoietic cell proliferation disorder related sequences and their complements, and as primers for the calated sequences and their complements, and as primers for the calated sequences. The nucleotide sequences from the present invention can subob a used for detecting a pracisopsition to, differentiation between suboliterative disorders. The present method enables allowing for improved and informed treatment of patients.
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disorders, comprises contacting a target nucleic acid with a reagent
that distinguishes between methylated and non-methylated CpG
                                                                                                                                                                                                                                                                                                         Berlin K, Braun A, Distler J, Guetig D, Howe A, Mueller J; Olek A, Piepenbrock C, Adorjan P, Grabs G, Lesche R, Leu B; Lewin A, Lipscher E, Maier S, Model P, Mueller V, Otto T; Pelet C, Schwope I, Ziebarth H;
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Pred. No. 2.5e+02;
0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 15; Page 44; 117pp; English.
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Matches 15; Conservative
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                                         Homo sapiens.
Synthetic.
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Human, haematopoietic cell proliferation disorder, cytostatic; gene therapy, lymphocytic leukaemia, acute myelogenous leukaemia; cytosine methylation state; probe; primer; ss.
                                03-OCT-2002
                                                              Berlin K,
                                                                 Olek A, I
Lewin A,
                  Synthetic
                                                                         Pelet
              Homo
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The present invention describes a method for detecting and differentiating between haematopoietic cell proliferative disorders associated with at least 1 gene and/or their regulatory regions in a subject. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least 1 reagent, which distinguishes between methylated and non-methylated cipG dinucleotides within the target nucleic acid. ABZ09861 to ABZ1118 represent specifically claimed nucleotide sequences from the present invention. Oligonucleotides from the present invention can be used: for differentiating between healthy haematopoietic cells and proliferative disorder haematopoietic cells; for differentiating between acute lymphocytic leukaemia and acute myelogenous leukaemia; as probes for determining the cytosine methylation state and/or single nucleotide polymorphisms (SNPs) of haematopoietic cell proliferation disorder related manning the nucleotide sequences from the present invention can also be used for detecting a predisposition to, differentiation between subclasses, diagnosis, prepagnosis, treatment and/or monitoring of haematopoietic cell proliferation ethod enables a highly specific classification of haematopoietic cell proliferative a highly specific classification of haematopoietic cell proliferative a highly specific classification of haematopoietic cell proliferative ö Detecting and differentiating between hematopoietic cell proliferative disorders, comprises contacting a target nucleic acid with a reagent that distinguishes between methylated and non-methylated CpG disorders allowing for improved and informed treatment of patients. Gaps , Braun A, Distler J, Guetig D, Howe A, Mueller J; Piepenbrock C, Adorjan P, Grabs G, Lesche R, Leu B; Lipscher E, Maier S, Model F, Mueller V, Otto T; Schwope I, Ziebarth H; .. Score 13.2; DB 1; Length 18; Pred. No. 2.5e+02; 0; Mismatches 3; Indels Sequence 18 BP; 1 A; 0 C; 8 G; 9 T; 0 other; Claim 15; Page 44; 117pp; English. 0 1.0%; 26-MAR-2002; 2002WO-EP03401. 26-MAR-2001; 2001US-278339. 15; Conservative (EPIG-) EPIGENOMICS AG. WPI; 2003-018942/01 WO200277272-A2 dinucleotides sapiens.

24 AACCAAACCCAGCTACGC 41 18 AACCAAACCAACCTACAC 1 Query Match Best Local Similarity Matches 15; Conserv

ABZ11021 standard; DNA; 18 RESULT 426 ABZ11021

(first entry) 16-JAN-2003 ABZ11021; 2X2X5X

Haematopoietic cell proliferation disorder related oligonucleotide #1161.

Human; haematopoietic cell proliferation disorder; cytostatic; gene therapy; lymphocytic leukaemia; acute myelogenous leukaemia; cytosine methylation state; probe; primer; ss.

Homo sapiens. Synthetic. WO200277272-A2.

03-OCT-2002

26-MAR-2002; 2002WO-EP03401.

26-MAR-2001; 2001US-27833P.

(EPIG-) EPIGENOMICS AG.

Mueller J; e R, Leu E; Berlin K, Braun A, Distler J, Guetig D, Howe A, Mueller J, Olek A, Piepenbrock C, Adorjan P, Grabs G, Lesche R, Leu E Lewin A, Lipscher E, Maier S, Model F, Mueller V, Otto T; Pelet C, Schwope I, Ziebarth H;

WPI; 2003-018942/01.

Detecting and differentiating between hematopoietic cell proliferative disorders, comprises contacting a target nucleic acid with a reagent that distinguishes between methylated and non-methylated CpG dinucleotides

Claim 15; Page 76; 117pp; English.

The present invention describes a method for detecting and differentiating between haematopoietic cell proliferative disorders associated with at least 1 gene and/or their regulatory regions in a subject. The method comprises contacting a target nucleic acid in a bological sample obtained from the subject with at least 1 reagent, which distinguishes between methylated and non-methylated CpG dinucleotides within the target nucleic acid ABZ09861 to ABZ1118 represent specifically claimed nucleotide sequences from the present invention. Oligonucleotides from the present invention can be used: for differentiating between healthy haematopoietic cells and proliferative disorder haematopoietic cells; for differentiating between acute myelogenous leukaemia; as probes for determining the cyrosine methylation state and/or single nucleotide polymorphisms (SNPs) of haematopoietic cell proliferation disorder cangilification of haematopoietic cell proliferation disorder related mys sequences and their complements; and as primers for the mys sequences and their complements from the present invention cangils be used for detecting a predisposition to, differentiation between to subclasses, diagnosis, prognosis, treatment and/or monitoring of haematopoietic cell proliferative disorders. The present method enables a highly specific classification of haematopoietic cell proliferative cell proliferative of angulations.

Seguence 18 BP; 8 A; 8 C; 1 G; 1 T; 0 other;

Gaps .. 1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 2.5e+02; ative 0; Mismatches 3; Indels Conservative Similarity Ma. Local S... 15; Query Match Matches

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41 AACCAAACCGACCTACAC 24 AACCAAACCCAGCTACGC

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ABZ11022 standard; DNA; 18 RESULT 427 ABZ11022 ID ABZ1 XX AC ABZ1 XX

ABZ11022;

Thu Jan

(first entry)

(first entry)

21-FEB-2002

ABC89050;

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Haematopoietic cell proliferation disorder related oligonucleotide #1162
                      Human; haematopoietic cell proliferation disorder; cytostatic; gene therapy; lymphocytic leukaemia; acute myelogenous leukaemia; cytosine methylation state; probe; primer; ss.
                                                                                       26-MAR-2002; 2002WO-EP03401
                                                                                                  26-MAR-2001; 2001US-278333P.
                                                                                                              (EPIG-) EPIGENOMICS AG.
                                                                                                                                                     WPI; 2003-018942/01.
                                                                WO200277272-A2.
                                               Homo sapiens.
16-JAN-2003
                                                                           03-OCT-2002
                                                    Synthetic.
                                                                                                                                     Lewin A,
Pelet C,
                                                                                                                         Berlin K,
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, Braun A, Distler J, Guetig D, Howe A, Mueller J; Piepenbrock C, Adorjan P, Grabs G, Lesche R, Leu B; Lipscher E, Maier S, Model F, Mueller V, Otto T; Schwope I, Ziebarth H;

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The present invention describes a method for detecting and differentiating between haematopoietic cell proliferative disorders associated with at least I gene and/or their regulatory regions in a subject. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least I reagent, which distinguishes between methylated and non-methylated CpG dinucleotides within the target nucleic acid. ABZ09861 to ABZ11118 represent specifically claimed nucleotide sequences from the present invention. Oligonucleotides from the present invention and be used: for differentiating between healthy haematopoietic cells and proliferative disorder haematopoietic cells, for differentiating between acute myelogenous leukaemia; as probes for determining the cytosine methylation state and/or single nucleotide polymorphisms (SNPs) of haematopoietic cell proliferation disorder related sequences and their complements; and as primers for the amplification of haematopoietic cell proliferation disorder related amplification of haematopoietic cell proliferation disorder related amplication disorder related amplication disorder related amplication disorde
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Detecting and differentiating between hematopoietic cell proliferative disorders, comprises contacting a target nucleic acid with a reagent that distinguishes between methylated and non-methylated CpG
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Sequence 18 BP; 9 A; 8 C; 0 G; 1 T; 0 other;
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                               0;
1.0%; Score 13.2; DB 1; Length 18; 13.3%; Pred. No. 2.5e+02; ve 0; Mismatches 3; Indel8
             83.3%;
                            15; Conservative
                Best Local Similarity
 Query Match
                              Matches
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18
24 AACCAAACCCAGCTACGC 41
                             AACCAAACCAACCTACAC
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ABC89050 standard; DNA; 13 BP. RESULT 428 ABC89050/c ID ABC85

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single nucleotide polymorphisms and cytosine methylation status
                                       Oligonucleotide SEQ ID NO 89067 for detecting SNP TSC0022361.
                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID 89067; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                              Berlin K;
                                                                                                                                                                06-APR-2001; 2001WO-IB00713.
                                                                                                                                                                                    07-APR-2000; 2000DE-1019173.
                                                                                                                                                                                                                              Piepenbrock C,
                                                                                                                                                                                                         (EPIG-) EPIGENOMICS AG.
                                                                                                                                                                                                                                                 WPI; 2001-657177/75.
                                                                                                                       WO200177384-A2.
                                                                                                     Homo sapiens
                                                                                                                                            18-OCT-2001
                                                                                                                                                                                                                              olek A,
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99899, ABH00010-ABH99989 and ABI00010-ABE9989, ABF00010-ABH99989 and ABI00010-ABH99989 and specification, but was obtained in electronic form part of the printed specification, but was obtained in electronic format from WIPO at fit. wipo.int/pub/published\_pot\_sequences.

Sequence 13 BP; 12 A; 0 C; 1 G; 0 U; 0 other;

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   Length 13;
                                 0; Indels
1.0%; Score 13; DB 1; Le
100.0%; Pred. No. 1.8e+02;
cive 0; Mismatches 0;
                  Local Similarity 100.
es 13; Conservative
      Query Match
                                      Matches
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Gaps

q à

(first entry) 21-FEB-2002 ABC89051;

BP.

ABC89051 standard; DNA; 13

ABC89051

Oligonucleotide SEQ ID NO 89068 for detecting SNP TSC0022361.

SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Homo sapiens. 

WO200177384-A2

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                     Oligonucleotide SEQ ID NO 100003 for detecting SNP TSC0024859
                                                                                                                                                                                                                                                                                               ABC99986 standard; DNA; 13 BP
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                                                        Olek A,
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single nucleotide polymorphisms and cytosine methylation status
                                                                               Claim 1; SEQ ID 100003; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                               Sequence 13 BP; 0 A; 2 C; 10 G; 1 T; 0 other;
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Best Local Similarity 100.0
....nhes 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE09989, ABF00010-ABE09989 and ABE00010-ABE09989 and shalloudous ABE180273 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single nucleotide polymorphisms and cytosine methylation status
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Pred. No. 1.8e+02;
0; Mismatches 0;
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                                                                                                            07-APR-2000; 2000DE-1019173
                                                                                                                                                                                            Piepenbrock C,
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.
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1.0%; Score 13; DB 1; Length 13;
100.0%; Pred. No. 1.8e+02;
tive 0; Mismatches 0; Indels
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(first entry)

Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE09989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABF9989 and SPECONDIO ABF9989 and ABF00010-ABF9989 and ABF00010-ABF9980 and ABF00010-ABF9980 and ABF00010-ABF9980 and ABF00010-ABF9980 and ABF9980 and ABF9980
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ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention.
NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pot_sequences.
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100.0%; Pred. No. 1.8e+02;
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABH00010-ABH99989 and ABL00110-ABC9989, abeceting cell type differentiation. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                 SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligoners for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligoners are also used for detecting cell type differentiation. ABC00010-ABE09989, ABF00010-ABF99989, ABH00010-ABH99989 and ABE10010-ABE182073 represent the oligoners described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at
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             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.
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100.0%; Pred. No. 1.8e+02;
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               This invention describes novel oligonuclectide primers or peptide nucleic acid (PMA) oligoners for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DMA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH00010-ABH9989 and ABC0010-ABF89989, ABF0010-ABH99989 and ABC0010-ABF80731 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                      ..
                                                                                                                                                                                                                                                                                                                                                                               1.0%; Score 13; DB 1; Length 13;
100.0%; Pred. No. 1.8e+02;
live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                  Sequence 13 BP; 3 A; 0 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                             ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
es 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
*555555555555
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ABH18097 standard; DNA; 13 RESULT 437 ABH18097/c

BP.

Oligonucleotide SEQ ID NO 218074 for detecting SNP TSC0053025. (first entry) 22-FEB-2002 ABH18097; 

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

18-OCT-2001,

06-APR-2001; 2001WO-IB00713.

07-APR-2000; 2000DE-1019173

(EPIG-) EPIGENOMICS AG

Olek A,

Piepenbrock C,

Berlin K;

WPI; 2001-657177/75.

set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single nucleotide polymorphisms and cytosine methylation status

triphosphates used as antiviral agents and synthons

Claim 179; Page 156; 259pp; English.

Claim 1; SEQ ID 218074; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a renge of diseases including immune system, gastrointestinal, respiratory, oligomers are also used for detecting cell type differentiation.

ABC00010-ABE09999, ABF00010-ABF99999, ABH00010-ABH99999 and ABC00010-ABE18073 represent the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed ffp.wipo.int/pub/published\_pct\_sequences.

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                     Gaps
                                                                                                                                                                                            Human, c-raf, A-raf, B-raf, hammerhead ribozyme, hairpin ribozyme, target, substrate, catalyst, modulation, expression, Raf gene, delivery, screening; identification, synthesis, deprotection, purification; carcias, non-hepatic ascites, infection; genetic drift, restenosis, rheumatoh, asthritis, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside
                     ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          i, Bellon L, Burgin A, Jarvis T;
, Matulic-Adamic J, McSwiggen JA;
Sweedler D, Thompson J, Workman CT;
Length 13;
                     0; Indels
                                                                                                                                                                          Human C-raf target sequence nucleotide position 2567.
Score 13; DB 1; Le
Pred. No. 1.8e+02;
                     Mismatches
1.0.
100.0%; Fig.
                                                                                                             AAV92063 standard; RNA; 14 BP.
                                                                                                                                                                                                                                                                                                                                        97US-0068212.
97US-0046059.
97US-004002.
97US-0056808.
97US-0061321.
97US-0061324.
                                                                                                                                                                                                                                                                                                                        98WO-US09249
                                        1184 TATAGGTGAGTGT 1196
                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                      (first entry)
                   13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Beigelman L,
                                                          13 raradgreagrer 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reynolds M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-009494/01.
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Karpeisky A,
                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                      18-FEB-1999
                                                                                                                                                                                                                                                                               WO9850530-A2
                                                                                                                                                                                                                                                                                                                        05-MAY-1998;
                                                                                                                                                                                                                                                                                                   12-NOV-1998.
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                                                                                                                                                                                                                                                                                                                                                                                    22-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Beaudry A,
                                                                                                                                  AAV92063;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Parry T,
                   Matches
                                                                                          RESULT 43
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A method has been developed for the identification of a nucleic acid capable of modulating a process in a biological system. The method comprises: (a) introducing into the system a random library of nucleic acid catalysts (NAC) having a substrate binding domain (BBD), comprising a random sequence, and a catalytic domain (CD); and (B) identifying NAC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules with endonuclease activity and catalytic activity, from the present invention, are used to modulate gene expression in plant and mammalian cells and to cleave target nucleic acid, particularly for treating systemic diseases caused by specific RNA, e.g. cancer, inflammation, psoriasis, non-hepatic ascites and infection. They may also be used to detect genetic drift and mutations in diseased cells and to determine

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28-NOV-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 440
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c-raf RNA. Specifically NACs with RNA-cleaving activity that modulate expression of the Raf gene, are used to treat cancer, restenosis, psoriasis or rheumatoid arthritis, or generally any condition associated with the level of c-raf. Introduction of sugar/phosphate modifications increases stability against nuclease and activity. AAV99922 to AAV93877 represent NACs that can be used in the method, specifically for modulating the expression of a Raf gene.
                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                           transplant rejection; rhemmatoid arthritis; psoriasis;
myocardial ischaemia; Kawasaki disease; septic shock; HIV;
human immunodeficiency virus; acquired immune deficiency syndrome;
                                                                                                                                                                                                                                                                        Human relA hammerhead ribozyme target sequence (nt. position 585).
                                                                                                                                                                                                                                                                                         Bnzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; chromic myelogenous leukaemia; CML; cancer; Philadelphia chromosome; inflammation; autoimmune disease; atheroselerosis; myocardial infarction; stroke; restenosis;
                                                                                                                   ô
                                                                                     Local Similarity 92.3%; Pred. No. 2e+02; es 12; Conservative 1; Mismatcher
                                                                             Sequence 14 BP; 3 A; 4 C; 5 G; 2 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-0380734.
94US-0201109.
94US-0227953.
94US-0227483.
94US-0227483.
94US-0227586.
94US-021932.
94US-0291433.
94US-0291433.
94US-0291433.
94US-0291433.
94US-0291433.
94US-0291433.
94US-0303039.
94US-0303039.
94US-0311749.
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94US-0334847.
94US-0337608.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95WO-IB00156
                                                                                                                                                                                                       AATS5016 standard; RNA; 15
                                                                                                                                                                                                                                             (updated)
(first entry)
                                                                                                                                      480 GGACTGCCGAGAC 492
                                                                                                                                                       1 GGACUGCCGAGAC 13
                                                                                                                                                                                                                                                                                                                                                                                                                               WO9523225-A2.
                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-NOV-1994;
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04-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-AUG-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-AUG-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-AUG-1994
                                                                                                                                                                                                                                             25-MAR-2003
                                                                                                                                                                                                                                                      18-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                            AIDS; ss.
                                                                                                                                                                                                                           AAT55016;
                                                                                                Query Match
                                                                                                                                                                                      RESULT 439
                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves relament of mRNA at the nucleotide base position indicated in the DB line.

The rela gene product is a subunit of the transcriptional regulator NF-kappab and is implicated specifically in the induction of inflammatory responses. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and spring structures and that contain potential hammerhead analysis. Ribozyme cleavage sites were identified by computer analysis. Ribozyme directed against these mRNA sequences were callysis and springed and synthesised with modifications that improve their nuclease resistance. The ribozymes are designed to cleave the target sequences and thereby inhibit relA expression, making them potentially useful for treating rheumatoid arthritis, restenosis to and asthrma as well as for increasing tolerance to transplanted tissues. The potential immunosuppressive properties of a ribozyme that cleaves relA mRNA means that uses are limited to local delivery, acute indications or ex vivo treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide array; genotyping; single base extension reaction; SBE; PCR primer; polymorphic locus; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                              Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
                                                                                                                                                                                                                               Dudycz LW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ribozymes having modified bases and methods for producing them for use in inhibiting disease related genes
                                                                                                                                                                                                                           Stinchcomb DT, Chowrira B, Direnzo A, Draper KG, Dudycz Grimm S, Karpeisky A, Kisich K, matulic-adamic J, Moswig Modak A, Pavco P, Beigleman L, Sullivan SM, Sweedler D; Thompson UD, Tracz D, Usman N, Wincott FE, Woolf T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.0%; Score 13; DB 1; Length 15; 92.3%; Pred. No. 2.1e+02; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Reverse primer #97 used in multiplexing PCR/SBE assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 4 A; 5 C; 5 G; 1 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; Page 228; 407pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0126473.
99US-0140359.
94US-0345516.
94US-0357577.
94US-0363233.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1066 CCCATCAGGCAGG 1078
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                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-FEB-2001 (first entry)
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Best Local Similarity 92.3
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1995-351090/45.
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                                                                              23-DEC-1994;
                                               L6-DEC-1994;
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Length 15;

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Human; phospholipid transfer protein; PLTP; SNP; atherosclerosis; single nucleotide polymorphism; high-density lipoprotein metabolism; allele-specific oligonucleotide; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                     Human phospholipid transfer protein gene ASO primer SEQ ID NO: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genotyping phospholipid transfer protein gene of individual for haplotyping individual's gene, comprises determining identity of nucleotide pair at polymorphic sites for two copies of PLTP gene present in the individual
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENA-) GENAISSANCE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-MAR-2000; 2000US-192127P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-MAR-2001; 2001WO-US08283
                                                                                                                                                                                                                                                                                                                    ABA81582 standard; DNA; 15
                                                                                                                                                                                                                                                         813 GCCGAGCGTCCTG 825
                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                        15 GCCGAGCGTCCTG 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Choi JY,
                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200172761-A2
                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                an J,
                        Ryder
                                                                                                                                                                                                                                                                                                    RESULT 441
                                                                                                                                                                                                                                        Matches
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BP.

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phospholipid transfer protein (PLTP) gene, involving determining the identity of the nucleotide present at one or more of the 25 polymorphic sites within the gene. This can be used to aid drug development for the treatment of diseases associated with different haplotypes of the PLTP gene, possibly including atherosclerosis. The present sequence is an allele-specific primer used for detecting polymorphisms in the PLTP gene
                                                                                                                                                                                                                                                            1.0%; Score 13; DB 1; Length 15;
86.7%; Pred. No. 2.1e+02;
tive 1; Mismatches 1; Indels
                                                                                                                                                                                                 Sequence 15 BP; 1 A; 7 C; 2 G; 4 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                             127 ACGGGACAGGGACGC 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IGF-I oligonucleotide #2661.
                                                                                                                                                                                                                                                                                                                                                                                                                       ASGGGATAGGGACGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF51701 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-MAR-2001
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           88888888888
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                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to an oligonucleotide array comprising oligonucleotide tags fixed to a solid substrate. The oligonucleotide tags fixed to a solid substrate. The oligonucleotide tarray is useful for genotyping a nucleic acid sample at one or more loci via single base extension (SEB) reactions. A pair of primers is used to amplify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNP). The present sequence is one of the primers used in the method of the present invention to amplify a polymorphic sample. The amplified nucleic acid product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to
                                                                                                                                                                                                          Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions
                                                                    5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                    Lockhart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .
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                                                                       Kaplan P, Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 BP; 2 A; 5 C; 6 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                 Example 7; Page 58; 83pp; English.
                                                                       Huang X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                form the oligonucleotide array.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13; Conservative
              (AFFY-) AFFYMETRIX INC.
                                                                          Hirschhorn JN,
                                                                                                                                                       WPI; 2000-656171/63
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Best Local Similarity
                                                                                                     Sklar P;
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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, piryriasis, ruba, pilaris, serborthoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other
                                                                                                                                 Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin discorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFB-2; IGFBP3; inflammation; psoriasis; pilatis; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neophasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neobascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Edmondson SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MURD-) MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 8; Page 78; 201pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-JUN-2000; 2000WO-AU00693.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wraight CJ, Werther GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-041421/05.
                                                                                                                                                                                                                                                                                                                                                                                 WO200078341-A1
                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                           28-DEC-2000.
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The present invention relates to a method for haplotyping the human

Claim 15; Page 13; 98pp; English.

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Gaps

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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide. (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein (IGFBP)-2 or IGFBB3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotide so of the present invention (see AAF45151 and AAF4513-1 The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition of warts, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic;
kidney disease, hyperproliferation of the inside of
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0
                                                                                                                                                                               Length 15;
                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                            Score 13; DB 1; Lo
Pred. No. 2.1e+02;
                                                                                                  Sequence 15 BP; 4 A; 2 C; 6 G; 3 T; 0 other;
                                                                                                                                                                                              100.0%; Pred. wv.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Edmondson SR;
          sclerotic disease, kidney disease, hype
blood vessels or any other hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MURD-) MURDOCH CHILDRENS RES INST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0140345,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IGF-I oligonucleotide #2662.
                                                                                                                                                                               1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF51702 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                       160 CGCTGATCCTCAA 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                Local Similarity 100.
Les 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF51702;
                                                                                                                                                                                       Query Match
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AAF51702/c

AC AAF517

AC ACCO

ACCO

AC ACCO

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Matches
              2000
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Sequence 15 BP; 4 A; 2 C; 6 G; 3 T; 0 other;

blood vessels or any other hyperplasia.

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The present invention relates to methods and oligonuclectides for forming a triple-helix comprising a double helical nucleic acid comprising first and second substantially complementary strands, and an oligonuclectide bound to a purine-rich target sequence within the double helical nucleic acid, where the oligonuclectide binds in a parallel and antiparallel orientation, respectively. to target sequences on alternate strands of the double helical nucleic acid. The method has therapeutic applications, where gene expression is controlled by selective triple-helix formation within expression is regulatory sequences of a target gene. The oligonucleotides can be used to form triple-helix sequences within genomic DNA for diagnostic and therapeutic purposes. The oligonucleotides can be selected to specifically bind to pathogenic double-stranded DNA including specific sequences required by pathogenicity and conviruence of the oligonucleotides can be selected to sequences required by pathogenicity. Alternatively, the oligonucleotide can be chosen to cauget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be chosen to carget a unique sequence of the oligonucleotide can be carged to carget a unique sequence of the carget and carget a unique sequence of the carget and carget a unique sequence of the carget and carget and carge
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A triple-helix comprising a double helical nucleic acid (DHNA) and an oligonuclectide which binds in parallel and antiparallel orientation, respectively, for targetting sequences on alternate strands of DHNA to control gene expression -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Triple-helix formation; purine-rich target sequence; double-helix DNA; gene expression; regulatory sequence; pathogenic double-stranded DNA; pathogenic bacteria; virus; replication; virulence; cancer; oncogene suppression; cancerous cell; cytostatic; antimicrobial; ss.
                                                   Gaps
                                                   0;
Length 15;
                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Triple helix forming associated oligonucleotide #36.
1.0%; Score 13; DB 1; Le
.00.0%; Pred. No. 2.1e+02;
                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CALY ) CALIFORNIA INST OF TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 6; Fig 20A; 108pp; English.
        1.0.0%; Pre-
                                                                                                                                                                                                                                                                                                       BP.
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                                                                                                                                                                                                                                                                                                       ABK98166 standard; DNA; 15
                                                                                                            160 CGCTGATCCTCAA 172
                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                        13; Conservative
                                                                                                                                                            14 cecrearcercan 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Beal PA;
     Query Match
Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-SEP-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                              07-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dervan PB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                             ABK98166;
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Gaps
                                                .,
                         Length 15;
                                              Indels
                                   2.1e+02;
                      1.0%; Score 13; DB 1;
BP; 0 A; 1 C; 0 G; 14 T; 0 other;
                                  100.0%; Pred. No. 2.1 cive 0; Mismatches
                                                                      1144 TTTTTTTTTT 1156
                                Local Similarity 100.
 Sequence 15
                      Query Match
                                             Matches
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rrrrrrcrrrr 13

RESULT 445

ABK98185 standard; DNA; 15 ABK98185;

BP.

(first entry) 07-OCT-2002

Triple helix forming associated oligonucleotide #49.

Triple-helix formation; purine-rich target sequence; double-helix DNA; gene expression; regulatory sequence; pathogenic double-stranded DNA; pathogenic bacteria; virus; replication; virulence; cancer; oncogene suppression; cancerous cell; cytostatic; antimicrobial; ss.

Synthetic.

US6403302-B1

11-JUN-2002

93US-0168920, 16-DEC-1993; 92US-0946976 17-SEP-1992; (CALY ) CALIFORNIA INST OF TECHNOLOGY.

Beal PA; Dervan PB,

WPI; 2002-536030/57.

A triple-helix comprising a double helical nucleic acid (DHNA) and an oligonucleotide which binds in parallel and antiparallel orientation, respectively, for targetting sequences on alternate strands of DHNA to control gene expression -

Example 7; Fig 24A; 108pp; English.

The present invention relates to methods and oligonucleotides for forming a triple-helix comprising a double helical nucleic acid comprising first and second substantially complementary strands, and an oligonucleotide bound to a purine-rich target sequence within the double helical nucleic acid, where the oligonucleotide binds in a parallel and antiparallel orientation, respectively, to target sequences on alternate strands of the double helical nucleic acid. The method has therapeutic applications, where gene expression is controlled by selective triple-helix formation within expression regulatory sequences of a target gene. The oligonucleotides can be used to form triple-helices, and are useful to detect the presence or absence of specific sequences within genomic DNA for diagnostic and therapeutic purposes. The oligonucleotides can be selected to specific sequences within genomic DNA for diagnostic and therapeutic purposes. The oligonucleotides can be selected to specific sequences within genomic DNA including specific specific and specifically bind to pathogenic double-stranded DNA including specific sequences required by pathogenic bacteria or viruses for replication or virule or viruses for replication or virulence, reducing their pathogenicity. Alternatively, the virulence, reducing their pathogenicity. Alternatively, the pligonucleotide can be chosen to target a unique sequence of the pathogen which is not found in the genome of pathogen's host. The oligonucleotides can be used in cancer treatment by way of triple-helix suppression of specific oncogenes including those of endogenous or viral origin. Such therapeutic oligonucleotides are capable of forming triple-helices with such sequences in cancerous cells containing the activated oncogene, so preferentially killing or repressing the cancer specific

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Gaps
causing cell. The present sequence represents an oligonucleotide used in the methods of the present invention.
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0
                                                                         Length 15;
                                                                                                     0; Indels
                                                                        Score 13; DB 1; Lo
Pred. No. 2.1e+02;
                                           Sequence 15 BP; 0 A; 1 C; 0 G; 14 T; 0 other;
                                                               1.0%; bcc.
100.0%; Pred. No. ...
                                                                                                     13; Conservative
                                                                     Query Match
Best Local Similarity
                                                                                                     Matches
 88333
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1144 TITITITICITITI 1156 rrrrrrrrrrr 13 Ч

à a RESULT 44 ABK95944/

ABK95944 standard; DNA; 15 BP

ABK95944;

(first entry) 24-SEP-2002

Human LIPE gene polymorphism detection ASO probe #6

Human; lipase; hormone sensitive; LIPE; isogene; obesity; probe; ss male sterility; polymorphism; allele-specific oligonucleotide; ASO.

Homo sapiens.

WO200240502-A2

23-MAY-2002.

16-NOV-2001; 2001WO-US43518.

16-NOV-2000; 2000US-249302P.

(GENA-) GENAISSANCE PHARM INC.

Rounds Koshy B, Chew A, Bentivegna SC, Anastasio AE,

WPI; 2002-519369/55.

Novel genetic variants of Lipase, Hormone-Sensitive isogenes, useful for improving efficiency and reliability in drug development for treating diseases associated with LIPE activity, e.g. obesity and male sterility

Claim 15; Page 14; 142pp; English.

The present invention relates to a new polynucleotide comprising a nucleotide sequence which comprises lipase, hormone sensitive (LIPE) isogenes. The invention is useful in screening for drugs targeting LIPE isogenes that are useful for treating obssity and male sterility. The isogenes that are useful for improving the efficiency and reliability of several steps in the discovery and development of drugs reliability of several steps in the discovery and development of drugs of its useful in studying the expression and function of LIPE, and in expressing LIPE protein for use in screening for candidate drugs to treat diseases related to LIPE activity. It is also useful in studying the effect of the variation on the biological activity of LIPE as well as on the binding affinity of candidate drugs targeting LIPE for the treatment of consists and male sterility. The invention is useful for studying the expression of LIPE isogenes in vivo, for in vivo screening and testing of the capteut against LIPE protein, and for testing the efficacy of the respective against LIPE protein, and for testing the efficacy of the appendic agents and compounds for treating obesity and male sterility in a biological system. The present nucleic acid sequence represents one the biology probes that were used in the invention to detect polymorphisms in the human LIPE gene. 

Sequence 15 BP; 5 A; 4 C; 1 G; 4 T; 1 other;

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The invention describes an isolated polynucleotide, which comprises genes and haplotypes of the small inducible cytokine subfamily A (Cys-Cys), member 20 (SCYA20) gene. The polynucleotide comprises polynorphic sites referred to as PS1-9 to designate the order in which they are located in the gene. The polymorphisms and haplotypes of SCYA20 gene are useful for validating whether SCYA20 is a suitable target for drugs to treat peoriasis and disorders associated with its abnormal expression or function, soreening for such drugs and reducing bias in clinical trials of such drugs. Haplotype information would be useful in improving the circlency and output of several steps in the drug discovery and development process, including target validation, identifying lead compounds, early phase clinical trials. The methods are useful in screening for compounds targeting SCYA20 to treat a specific condition criseases predicted to be associated with SCYA20 activity, e.g. psoriaeis. This sequence represents an allele specific oligonucleotide (ASO) probe used to identify polymorphisms in the SCYA20 gene.
                                         ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New genetic variants comprising haplotypes of the small inducible cytokine subfamily A, member 20 (SCYA20) gene, useful in improving the efficiency drug screening protocols for compounds (e.g. antipsoriatic drug) targeting SCYA20
                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                         Small inducible cytokine subfamily A (Cys-Cys) member 20; SCYA20; polymorphism; haplotype; psoriasis; gene expression; ASO; allele specific oligonucleotide; probe; ss.
                                         .
0
Query Match 1.0%; Score 13; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 2.1e+02; Matches 13; Conservative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                    SCYA20 allele specific oligonucleotide probe #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sausker EA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 2 A; 7 C; 1 G; 4 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Russo DP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 14; Page 13; 62pp; English.
                                                                                                                                                                                                                    ABK81414 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GENA-) GENAISSANCE PHARM INC
                                                                                440 GAAAGTTGCTGAAGT 454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-OCT-2001; 2001WO-US46093.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-OCT-2000; 2000US-241725P.
                                                                                                   15 GATAGTTSCTGAAGT 1
                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bieglecki KM, Chew A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-435525/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200232927-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                                                                                                                                                                                              13-AUG-2002
                                                                                                                                                                                                                                                         ABK81414;
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(GENA-) GENAISSANCE PHARM INC.

Kliem SE,

Duda A,

13-SEP-2000; 2000US-232468P.

13-SEP-2001; 2001WO-US28780.

WO200222650-A2.

21-MAR-2002

/\*tag= .a /note= "polymorphic site indicated by an ambiguity base"

Location/Qualifiers

misc\_feature

Homo sapiens

Human; period (Drosophila) homologue 1; PER1; polymorphic variant; polymorphic site; genotyping; haplotyping; circadian rhythm regulation; single nucleotide polymorphism; SNP; gene; probe; ss.

Human PER1 allele specific oligonucleotide probe SEQ ID NO:24.

(first entry)

12-JUL-2002

ABL52099;

ABL52099 standard; DNA; 15 BP.

ABL52099

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The present invention describes an isolated human period (Drosophila)
homologue 1, (PER1) polynucleotide (I) comprising a sequence which is a
polymorphic variant for a reference sequence (RAELSTOT) for the PER1 gene
to till fragment, or a polymorphic variant of a reference sequence
(ABESZOTS) for a PER1 CDNA or its fragment. The present invention also
describes methods for genotyping and haplotyping the PER1 gene of an
individual. (I) is useful in studying the expression and function of
PER1, and in expressing PER1 protein for use in screening for candidate
crherapeutic purposes. A recombinant non-human organism transformed or
transfected with (I) can be used for studying expression of the PER1
isogenes in vivo, for in vivo screening and testing of drugs targeted
against PER1 protein, and for testing the efficacy of therapeutic agents
and compounds for disorders associated with circadian rhythm regulation.
The present sequence represents an allele specific oligonucleotide probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel isolated human period Drosophila homolog 1 polynucleotide, useful for therapeutic purposes, for studying the expression and function of the polynucleotide, and for expressing the homolog -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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86.7%; Pred. No. 2.1e+02;
ttive 1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 2 A; 5 C; 4 G; 3 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 17; Page 14; 162pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           287 CAGCAGCAATGTCTG 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAGCAGCRCTGTCTG 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-393941/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention.
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ABK67885/c RESULT 449

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Gaps

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1.0%; Score 13; DB 1; Length 15; 86.7%; Pred. No. 2.1e+02; ative 1; Mismatches 1; Indels

1199 GACCTTCACACCTCC 1213

à g

13; Conservative

Query Match Best Local Similarity Matches 13; Conserva

1 GACCTTCRCACCTTC 15

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The invention relates to single nucleotide polymorphisms in the gene encoding human alcohol dehydrogenase 7 (class IV), mu or sigma polypeptide (ADH7). A method for haplotyping the ADH7 gene in an individual comprises identifying the nucleotide at one or more individual comprises identifying the nucleotide at one or more polymorphic sites and determining whether one of the copies of the gene is defined by one of the ADH7 haplotypes given in the specification or whether both copies are defined by a haplotype pairs can be useful in genotyphing, whereby all possible haplotype pairs can be assigned to specific genotypes. An association between a trait and a haplotype or haplotype pair of the ADH7 gene can be identified by comparing the frequency of the haplotype pair in a reference population, where a higher haplotype or haplotype pair in a reference population, where a higher haplotype or haplotype pair. ADH7 and its corresponding DNA are used for studying the expression and function of ADH7, and in screening for candidate drugs to treat diseases related to ADH7 activity, such as concerned pair allele-specific oligonucleotide probes used for detecting ADH7 gene
                                                                                                                                                    Human; alcohol dehydrogenase 7 class IV; ADH7; probe; ss; haplotype pair; haplotyping; cytostatic; antiparkinsonian; sigma polypeptide; cancer; single nucleotide polymorphism; genotyping; gene therapy; drug screening; mu polypeptide; Parkinson's disease; chromosome 4q23-q24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New genetic variants of the alcohol dehydrogenase 7 (class IV) mu or sigma polypeptide (ADH7) gene having polymorphisms, useful for treating disorders affected by ADH7 isogene expression or function, e.g. cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sausker EA;
                                                                                                                   Human ADH7 gene allele-specific oligonucleotide probe #12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.0%; Score 13; DB 1; Length 15; 86.7%; Pred. No. 2.1e+02; Live 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Parks KE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Koshy B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 BP; 2 A; 3 C; 6 G; 3 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kazemi A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 16; Page 13; 113pp; English.
ABK67885 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GENA-) GENAISSANCE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                               19-SEP-2001; 2001WO-US29508
                                                                                                                                                                                                                                                                                                                                                                                                                    19-SEP-2000; 2000US-233520P
                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 86.77
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Finkel K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or Parkinson's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-352009/38.
                                                                                                                                                                                                                                                                                                 WO200224958-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bieglecki KM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymorphisms
                                                                                                                                                                                                                                                           Homo sapiens.
                                                                             02-JUL-2002
                                     ABK67885
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Human, B-factor; properdin, BF; primer; ss; gene therapy; drug screening;
antidiabetic; dermatological; diabetes; immunosuppressive;
antiinflammatory; systemic lupus erythematosus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to oligonucleotide analogues comprising a protein nucleic acid molecule (PNA) monomer. They are used in the detection and separation of nucleic acid molecules and as probes, primers, linkers, adapters and antisense agente on solid supports. Modifications enhance their use as capture and detection probes e.g. by the incorporation of biotin, digoxigenen, radioisotopes, fluorescent labels such as 1 knorescent and reporter molecules such as alkaline phosphatase. They are also used for enhancing or inhibiting the activity of an enzyme or cellular activity. The compounds are stable to nucleases and proteases, have high affinity, binding specificity and solubility. The polyamide backbone of PNAs is resistant to both nucleases and proteases. PNAs bind mucleic acid molecules with greater affinity than DNA or RNA concentration. The compounds are relatively simple to synthesize and are used in a wide variety of applications. This sequence
                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonuclectides analogues useful in detection, separation and purification of nucleic acid molecules, comprise monomers, dimers and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene allele-specific oligonucleotide sequencing primer #30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              represents a DNA oligomer which is used to represent the effect of
                                              Nucleotide sequence of oligomer # 12 used to compare mismatches.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .
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                                                                                                                                                                                                                                                                                                                                           Archdeacon D, Archdeacon tova A, Choob M, Hondorp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13; DB 1; Le
Pred. No. 2.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 0 A; 1 C; 0 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              single base mismatches on oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                 Protein nucleic acid molecule; PNA; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 20; Page 123; 197pp; English.
                                                                                                                                                                                                                                                                                                                                          Efimov V, Fernandez J, Archdead
Chakhmakhcheau O, Buryakova A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0°,
100.0%;
10°,0%;
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                                                                                                                                                                                                                                                     14-MAR-2000; 2000US-189190P.
                                                                                                                                                                                                                    13-MAR-2001; 2001WO-US08111.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1144 TITITITITI 1156
              18-JUN-2002 (first entry)
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                                                                                                                                                                                                                                                                                                         (ACTI-) ACTIVE MOTIF.
                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-041177/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                   WO200168673-A1
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                                                                                                                                                                                      20-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               oligomers -
                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK64023;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human BF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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ABK64023
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Gaps

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1052 ACAGCCCTGGCCTTC 1066

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15 ACAGCCAYGGCCTTC

ABA97405 standard; DNA; 15 BP.

RESULT 450 ABA97405 ABA97405;

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The invention relates to single nucleotide polymorphisms in the gene encoding the human B-factor properdin protein (BF). A method for haplotyping the BF gene in an individual comprises identifying the nucleotide at one or more polymorphic sites and determining whether one of the copies of the gene is defined by one of the BF haplotypes given in the specification or whether both copies are defined by a haplotype pair. This method is useful in genotypes. An association between a trait and a haplotype or haplotype or haplotype pair of the BF gene can be identified by comparing the frequency of the haplotype pair in a population exhibiting the trait with the frequency of the haplotype or haplotype pair in a reference population, where a higher haplotype or haplotype or haplotype pair in a reference population, where a higher haplotype or haplotype pair in a reference population indicates the trait is associated with the haplotype or haplotype pair. Br and its corresponding DNA are used for studying the expression and function of BF, for use in screening for candidate drugs to treat diseases related to BF activity, such as diabetes and systemic lupus erythematosus. Sequences ABK63994-ABK64049 or represent allele-specific sequencing primers used to detect human BF gene
                                                                                                                                                                                                                                                                                                 New genetic variants having polymorphisms in the B-Factor, Properdin (BF) gene, useful for studying the function of BF, and for treating disorders affected by expression or function of the BF isogene -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; single nucleotide polymorphism; SNP; GHRHR; chromosome 7p14; growth hormone releasing hormone receptor; haplotyping; genotyping; isolated growth hormone defliciency; IGHD; pituitary adenoma; ASO; allele-specific oligonucleotide; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.0%; Score 13; DB 1; Length 15; 86.7%; Pred. No. 2.1e+02; 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ASO primer #7 to detect human GHRHR gene polymorphisms.
                                                                                                                                                                                                                              Koshy B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 BP; 3 A; 3 C; 8 G; 0 U; 1 other;
                                                                                                                                                                                                                              Kazemi A,
                                                                                                                                                                                                                                                                                                                                                                             Claim 17; Page 16; 151pp; English.
                                                                                                                                                                                      (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAS19628 standard; DNA; 15 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               129 GGGACAGGGACGCCC 143
                                                                                                             29-AUG-2001; 2001WO-US27098
                                                                                                                                                29-AUG-2000; 2000US-228940P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GGGACAGGGAGGCYC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                          Anastasio AE, Finkel K,
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                                                                                                                                                                                                                                                               WPI; 2002-304244/34.
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Best Local Similarity
                                 WO200218414-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200179239-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymorphisms.
 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-MAR-2002
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                                                                       07-MAR-2002
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The present invention relates to novel single nucleotide polymorphisms (SNPs) in the human growth hormone releasing hormone receptor (GHRHR) gene located on chromosome 7pl4, and methods for haplotyping and/or genotyping the GHRHR gene. The methods of the invention make use of allele-specific oligonucleotides (ASOS) as probes and primers and/or primer-extension oligonucleotides for detecting the GHRHR gene polymorphisms. The polymorphisms are useful for the treatment of diseases associated with GHRHR activity, such as isolated growth hormone deficiency (IGHD) and pituitary adenomas.

AASI9622-AASI9647 represent ASO primers for detecting human GHRHR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                   Genotyping human Growth hormone releasing hormone receptor gene of individual for determining haplotype of individual by determining identity of nucleotide pair at specific polymorphic sites for two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Colony stimulating factor 1 receptor; CSF1R; polymorphic variant; cytostatic; gene therapy; malignant histiocytosia; isogene; myeloid malignancy; inflammatory disorder; transgenic animal; haplotype; genotype; human; allele specific oligonucleotide; ASO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Colony stimulating factor 1 receptor (CSFIR) oligonucleotide #33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                          Sausker EA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.0%; Score 13; DB 1; Length 15;
86.7%; Pred. No. 2.1e+02;
ative 1; Mismatches 1; Indels
                                                                                                          Denton RR, Nandabalan K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 7 A; 2 C; 5 G; 0 U; 1 other;
                                                                                                                                                                                                                                             Claim 16; Page 14; 90pp; English.
                                                                           (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1158 GAAGTAAAGCAGCTA 1172
              17-APR-2001; 2001WO-US12453.
                                          17-APR-2000; 2000US-197978P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GAAGGAAAGCAGCYA 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Koshy B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAS98667/c
ID AAS98667 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAR-2002 (first entry)
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                                                                                                                                      WPI; 2002-066342/09.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           gene polymorphisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
les 13; Conserv
                                                                                                        Choi JY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           choi JY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200179225-A2.
                                                                                                                                                                                                                    copies of gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                        Chew A,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 453
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Novel polymorphic variants of colony stimulating factor 1 receptor useful in studying expression and function of the protein, useful for screening candidate drugs to treat diseases e.g. inflammatory disorders

15; 164pp; English. Page

The invention describes a novel isolated polymucleotide (I) comprising a sequence which is a polymorphic variant (PV) of a reference sequence for colony stimularing factor I receptor (CSFIR) gane, found on The colony stimularing factor I receptor (CSFIR) gane, found on The polypeptide are useful for improving the discovery and development of anilarmmatory discovery and the haplotypes can be used to validate CSFIR activity, e.g., and inflammatory discorders and the haplotypes can be used to validate CSFIR a candidate target for treating a specific condition or disease predicted to be associated with CSFIR and in the validate CSFIR species of an individual can also be used in studying the expression and therapeutic treatments. (I) is useful in studying the expression and function of CSFIR, protein for use in screening for candidate drugs to treat diseases related to CSFIR argeting CSFIR, Antibodies are useful in a variety of diagnostic and prognostic formats and therapeutic useful in a variety of diagnostic and prognostic formats and therapeutic contents in vivo screening and testing of drugs trargeted against CSFIR protein, and for testing the efficacy of therapeutic agents and compounds. Allele specific oligonucleotides (ASO) are useful as probes and primers, and for assaying a polymorphism in the target region. Without requiring any a priori knowledge of the phenotypic effect of any particular CSFIR or are useful as probes and primers, and for assaying a polymorphism in the efficacy in clinical trials. This sequence is an allele specific oligonucleotide probe used for detecting CSFIR gene polymorphisms, coligonucleotide probe used for detecting CSFIR gene polymorphisms, coligo

Sequence 15 BP; 2 A; 4 C; 4 G; 4 T; 1 other;

Gaps ;0 1.0%; Score 13; DB 1; Length 15; 36.7%; Pred. No. 2.1e+02; ive 1; Mismatches 1; Indels ilarity 86.7%; Conservative Local Similarity nes 13; Conserv Query Match Best Loca Matches

812 AGCCGAGCGTCCTGA 826

à

15 AGCCAAGYGTCCTGA 1 qq

AAS94594/c ID AAS94594 standard; DNA; 15 BP. RESULT 454

AAS94594;

Human PLTP gene allele-specific oligonucleotide sequencing primer #3.

(first entry)

14-FEB-2002

Human; phospholipid transfer protein; PLTP; haplotyping; haplotype pair; single nucleotide polymorphism; genotyping; gene therapy; drug screening; binding affinity; atherosclerosis; ss; sequencing primer; PCR primer;

Homo sapiens.

WO200172966-A2.

04-OCT-2001

26-MAR-2001; 2001WO-US09776.

24-MAR-2000; 2000US-192127P. 

(GENA-) GENAISSANCE PHARM INC

Koshy B; Choi JY, Chew A,

WPI; 2002-010724/01.

New isolated polynucleotide which is polymorphic variant of phospholipid transfer protein (PLTP) gene, having any one of polymorphic sites PSI-PS25, for studying function of PLTP, and expressing PLTP protein

Claim 15; Page 72; 99pp; English.

The invention relates to single nucleotide polymorphisms in the gene encoding the human phospholipid transfer protein (PLTP). A method for haplotyping the PLTP gene in an individual comprises identifying the nucleotide at one or more polymorphic sites and determining whether one of the copies of the gene is defined by one of the PLTP haplotypes given in the specification or whether both copies are defined by a haplotype pair. This method is useful in genotypies. An association between a crait and a haplotype or haplotype pair of the haplotype or haplotype pair of the haplotype or haplotype pair of the haplotype or haplotype or haplotype pair of the haplotype or haplotype pair in a population exhibiting the trait with the frequency of the haplotype or haplotype pair. In a reference population, where a higher haplotype or haplotype or haplotype pair in a reference population of PLTP, for use in screening frequency in the trait population indicates the trait is associated with the haplotype or haplotype pair. PLTP and its corresponding DNA are used for studying the expression and function of PLTP, for use in screening for studying the effect of variation on the sequences are also useful for studying the effect of variation on the candidate drugs targeting PLTP for treating atherosclerosis. Sequences AASS4691 represent allele-specific oligonucleotide probes, and purpose. polymorphisms. NAME OF THE PART O

Sequence 15 BP; 1 A; 7 C; 2 G; 4 T; 1 other;

Gaps .. Score 13; DB 1; Length 15; Pred. No. 2.1e+02; Indels 1.0%; 86.7%; Best Local Similarity 86.73 Matches 13; Conservative Query Match

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455 RESULT 45 ABL42625

ABL42625 standard; DNA; 15

BP.

ABL42625;

(first entry) 11-APR-2002

Hairpin beacon target hybridisation oligonucleotide #4.

Hybridisation, thermodynamic, computer readable storage medium, probe; target, molecular beacon, duplex; hairpin, ss. 

Synthetic.

WO200194611-A2

13-DEC-2001.

07-JUN-2001; 2001WO-US18424

07-JUN-2000; 2000US-209778P

(UYWA-) UNIV WAYNE STATE

Santalucia J, Peyret N;

WPI; 2002-122125/16.

us09904568-3.rng

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The present invention describes a method for predicting nucleic acid hybridisation thermodynamics (HT) comprising providing a database of thermodynamic parameters (FT), receiving hybridisation information which represents a sequence, receiving correction data, and a first set of data which represents hybridisation conditions, and calcularing HT correction data and the first set of data. Also described are: (1) a computer-readable storage medium having stored in it, a database of TP and a computer program which executes the above method; and (2) a system computer program which executes the above method; and (2) a system and a computer program which executes the above method; and (2) a system computer program which computers at least one for predicting nucleic acid HT, comprising a database of TP, units for receiving hybridisation information which represents at least one sequence and for receiving correction data, receiving a first set of data which represents hybridisation conditions and unit for calculating HT. The method and system are useful to optimise and predict probe-target hybridisation. The method and system takes into account of single strand thermodynamics not taken into account by prior art methods. ABL42498 to ABL42626 represent oligonucleotide sequences which are used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
            Predicting nucleic acid hybridization thermodynamics based on hybridization information, thermodynamic parameter, correction data and first set of data which represents hybridization conditions -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide #4 able to covalently cross-link to target DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.0%; Score 13; DB 1; Length 15; 100.0%; Pred. No. 2.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       deoxyribonucleic acid; major groove; ethanoamino group; aziridinylcytosine; cross-linking group; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 BP; 0 A; 1 C; 4 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  note= "N4N4-ethanocytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Prea. xv.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                   Disclosure; Fig 8; 100pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= b
/mod_base= m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ20008 standard; DNA; 16 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1146 TTTTTCTTTTGG 1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91WO-1003680
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 TTTTTCTTTTGG 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GILE-) GILEAD SCIE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAY-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-APR-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ20008;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Gaps
                                                                                                                              oligo is one of four oligonucleotides which were designed to specifically bind and cross-link to the duplex target sequence AAQ20004. Oligo #4 with its internal cross-linking group was less effective than the other oligonucleotides with terminal
                                                                                                                    The 3' end of this oligonucleotide carries 1,3-propanediol. The
                                                                                                                                                                                                                                                       .
0
                                             New sequence-specific non-photo-activated crosslinking agents -
bind to the major groove of duplex DNA and are esp. useful for
treating latent infections e.g. HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide #2 able to covalently cross-link to target DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New sequence-specific non-photo-activated crosslinking agents -
bind to the major groove of duplex DNA and are esp. useful for
treating latent infections e.g. HIV
                                                                                                                                                                                                                               Length 16;
                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                           deoxyribonucleic acid; major groove; ethanoamino group; aziridinylcytosine; cross-linking group; ss.
                                                                                                                                                                                                                              Score 13; DB 1; Le
Pred. No. 2.3e+02;
                                                                                                                                                                                                      Seguence 16 BP; 0 A; 2 C; 0 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note== "N4N4-ethanocytosine"
                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                               cross-linking groups. See also AAQ20005-7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                    1.0%; Scor
100.0%; Pred
                                                                                              Example 2; Page 21; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         base= OTHER
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/mod_base= m5c
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                                                                                                                                                                                                                                                                                                                                                              AAQ20006 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                            1144 TTTTTTTTTT 1156
Matteucci MD, Krawczyk S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Krawczyk S;
                                                                                                                                                                                                                                                                                                                                                                                                           01-APR-1992 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               *tag= a
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                                                                                                                                                                                                                                                                                                    1 Trititicitit 13
                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GILE-) GILEAD SCIE INC.
                       WPI; 1992-007480/01.
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                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified base
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                                                                                                                                                                                                                                                                                                                                                                                     AAQ20006;
                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                       RESULT 457
                                                                                                                                                                                                                                                      Matches
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Length 17;

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The 3' end of this oligonucleotide carries 1,3-propanediol. The oligo is one of four oligonucleotides which were designed to specifically bind and cross-link to the duplex target sequence AAQ20004. Oligo #2 has the covalent cross-linking group, i.e. N4N4-ethanocytosine, at its 3' end. An assay for crosslinked triple helix showed considerable reaction with Oligo #2 and with Oligo #1 (see AAQ20005) which has the crosslinking group at the 5' end. The most complete reaction was seen with Oligo #3 (see AAQ2007) having N4N4-ethanocytosine at both the 5' and 3' termini, A control oligo with no cross-linking group showed no reaction. The half-life of the cross-linking reaction for Oligo #2 was ca. 1 hr (1 microM);
                                                                                                                                                        Oligo #1 showed a rate four times slower. See also AAQ20008
                                                                                                                                                                              Sequence 17 BP; 0 A; 3 C; 0 G; 14 T; 0 other;
        Example 2; Page 20; 42pp; English.
                                                                                                                                                                                                Query Match
Best Local Similarity 100.
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pure and
                                                                                                                                                                                                                                                                                                                                                      AAQ33416;
                                                                                                                                                                                                                                                                                                        RESULT 458
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequences given in AAQ33414-17 are probes which were used to recognise polymorphic sites within the full length acid sphingomyelines (ASM) gene. This was done to determine the population frequency of the different alleles. The template DNA was genomic DNA from normal Caucasian individuals. Certain mutations in the ASM gene lie. R496L, deltaR608 and L302 have been found to correlate with Neimann-Pick disease (NPD). See also AAQ33390-423.
                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    s and recombinant acid sphingomyelinase and its nucleic acid treatment and diagnosis of Niemann-Pick disease
                                                                   ö
1.0%; Score 13; DB 1; Length 17;
100.0%; Pred. No. 2.5e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic site recognition probe, Arg-506.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 12; 50pp; English
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                                                                                                                                                                                                                                                                                                                                  AAQ33416 standard; cDNA; 17 BP
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                                                                                                                              1144 TTTTTTTTTTT 1156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schuchman EH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                  rrrrrrcrrrr 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                       (updated)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-MAY-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
06-MAY-1993
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Sequence 17 BP; 4 A; 6 C; 3 G; 4 T; 0 other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                               Gaps
                                                                                                                                                                                                                                                                                                          Primer; polymerase chain reaction; amplify; mutant; K-ras; PCR;
flanking region; amplification; probe; detection; sputum; diagnosis;
benign; malignant; neoplasm; lung; lung cancer; head; neck; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting target nucleic acid in mammalian sputum - particularly for diagnosis of lung neoplasia involving mutation(s) in the K\text{-}\mathrm{ras}
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Pred. No. 2.5e+02;
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                               Indels
1.0%; Score 13; DB 1; Le 100.0%; Pred. No. 2.5e+02; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 3 A; 9 C; 3 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Prea. ...
                                                                                                                                                                                                                                                                           p53 detection probe, (codon 178 ins 1 C).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYJO ) UNIV JOHNS HOPKINS SCHOOL MED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 36; 122pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oncogene or p53 tumour suppressor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT95093 standard; DNA; 17 BP.
                                                                                                                                                                               BP.
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                                                                                                                                                                               AAQ92208 standard; DNA; 17
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                                                               631 CTCCAGGAGCTCT 643
                                                                                                                                                                                                                                                 (first entry)
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Best Local Similarity 100.
Matches 13; Conservative
                                                                                              Crccaddadcrcr 17
 Query Match 1.03
Best Local Similarity 100.
Matches 13; Conservative
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                                                                                                                                                                                                                                                 12-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-NOV-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sidransky D;
                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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                                                                                                                                                                                                                  AAQ92208;
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The present sequence is a probe for a human acid sphingomyelinase (ASM) genomic DNA codon 506 mutation.

C Diagnosing Type A or B Niemann-Plck disease (NPD), or identifying a person as having the potential to genetically transmit Type A or B NPD, comprises detecting a recessive mutation in the ASM gene, which results in an alteration of at least 1 amino acid in the ASM amino acid sequence. The method is especially useful for prenatal diagnosis in Ashkenazi Jewish populations. The mutation is a Arg496Leu, deltaArg608, Leu302Pro or fsp330, where fsp330 is a crame shift mutation comprising a cytoshen deletion in ASM codon containing portions of the ASM gene by PCR using primers containing portions of the ASM gene by PCR using primers and sequencing the amplified DNA or subjecting it to a mid sequence, and sequence encoding it can also be used in the apprencing or the CDNA sequence encoding it can also be used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Diagnosing Type A or B Niemann-Pick disease - by detecting recessive mutation in acid sphingomyelinase gene
                                                                            for acid sphingomyelinase genomic DNA codon 506 mutation.
                                                                                                         Prenatal diagnosis; Type A; Type B; Niemann-Pick disease; identification; potential genetic transmiter; detection; recessive mutation; acid sphingomyelinase; Ashkenazi Jew; human; treatment; codon 506 mutation; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human EGF-R target sequence nucleotide position 73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 4 A; 6 C; 3 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Columns 51-52; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                         (MOUN ) MOUNT SINAI SCHOOL MEDICINE.
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Query Match

Best Local Similarity

Local 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                          Schuchman EH;
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                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-558133/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      treatment of NPD.
                                               17-FEB-1998
                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                           27-MAY-1994;
                                                                                                                                                                                                                                                                                                                                         27-MAY-1994;
03-MAY-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                          Desnick RJ,
                                                                                                                                                                                                                                           US5686240-A.
                                                                                                                                                                                                                                                                            11-NOV-1997.
                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV97224;
               AAT95093;
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                                                                                Probe
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ID AAV9'
XX AAV9'
XX DT 17-MJ
XX DE Humal
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94US-0250740. 91US-0695572. 94US-0250740.

The present invention describes enzymatic nucleic acid molecules (NAMS) which specifically cleave RNA derived from an epidermal growth factor receptor (EGF-R) gene. AAV97221 to AAV98043 and AAV98979 to AAV99090 represent specifically claimed target sequence from human EGF-R. AAV98044 haty98866 and AAV98867 to V9878 represent hammerhead ribozymes and hairpin ribozymes respectively for human EGF-R. The NAMS are useful for cleaving EGF-R RNA in the treatment of a condition associated with EGFR expression levels e.g. to inhibit cell proliferation in the prevention or treatment of cancers. The NAMS can also be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of EGF-R RNA in a cell. Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss. Human, epidermal growth factor receptor; BGFR, BGF-R, target sequence, hammerhead ribozyme, hairpin ribozyme; inhibition; cell proliferation; cancer; genetic drift; detection; mutation; ss. Enzymatic nucleic acids - which cleave RNA derived from an epidermal growth factor receptor, useful for inhibiting cell proliferation and for treating cancers Gaps ö Length 17; 0; Indels 1.0%; Score 13; DB 1; Le 100.0%; Pred. No. 2.5e+02; Human HTPL scanning oligonucleotide SEQ ID 381. Sequence 17 BP; 1 A; 8 C; 5 G; 3 U; 0 other; 100.0%; Pred. No. ... McSwiggen JA; Claim 5; Page 68; 109pp; English. BP. 97US-0985162. 97US-0036476. 98WO-US00730. ABV79135 standard; DNA; 17 200 CGGACGCCGACGA 212 (RIBO-) RIBOZYME PHARM INC. (first entry) Local Similarity 100. nes 13; Conservative Ŋ 17 cggacgccgacga WPI; 1998-437449/37 (UYAS-) UNIV ASTON. Akhtar S, Fell P, EP1229046-A2 04-DEC-1997; 31-JAN-1997; Homo sapiens 14-JAN-1998; WO9833893-A2 03-JAN-2003 07-AUG-2002. 06-AUG-1998 ABV79135; Query Match RESULT 462 Matches Homo ABV79135 g à

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Gaps

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Length 17; 0; Indels

Score 13; DB 1; Le Pred. No. 2.5e+02;

1.0%; 2

Mismatches

0;

07-AUG-2002

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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78752 and ABB98519 to ABB98520). HTPL broates in (HTPL, see ABV78759 to ABV78752 and ABB98519 to ABB98520). HTPL two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is contacted to the shared structure of a medicament, and the HTPL gene was mapped to human chromosome 1012.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of the HTPL. Such disorders include disorders of testis, or adrenal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, sheletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligomucleotide was used in an example from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                        Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 113; 718pp; English.
                                          30-JAN-2001; 2001WO-US00663.
30-JAN-2001; 2001WO-US00664.
30-JAN-2001; 2001WO-US00665.
30-JAN-2001; 2001WO-US00667.
30-JAN-2001; 2001WO-US00668.
30-JAN-2001; 2001WO-US00688.
                                                                                                                                                                                                                09-OCT-2001; 2001US-0327898
2002EP-0001167
                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-676582/73.
                                                                                                                                                                                                                                                              (AEOM-) AEOMICA INC
28-JAN-2002;
                                                                                                                                                                                                                                                                                                             Zhan J;
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0
                              Gaps
                             ;
0
Length 17;
1.0%; Score 13; DB 1; Length 17;
100.0%; Pred. No. 2.5e+02;
ive 0; Mismatches 0; Indels
             Local Similarity 100.
les 13; Conservative
Query Match
                              Matches
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ABV79136 standard; DNA; 17 BP. RESULT 463 

ABV79136;

03-JAN-2003 (first entry)

Human HTPL scanning oligonucleotide SEQ ID 382.

Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis syrpessed Patched like protain; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal musole; colon; male infertility; cancer; ss.

Homo sapiens.

EP1229046-A2

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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL brotein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL to be two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-8 (S for short) compared to HTPL-10 (L for long). HTPL shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The shares atructural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was important in regulating male germ cell development, and the HTPL, and in therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of the HTPL. Such disorders include disorders expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and forcal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonuclectide was used in an example from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 3 A; 6 C; 7 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; Page 113; 718pp; English.
                                                                                                                                                                                               30-JAN-2001; 2001WO-US00669.
23-MAY-2001; 2001US-0864761.
09-OCT-2001; 2001US-0327898.
                                                                                                                                                      2001WO-US00667.
                                             28-JAN-2002; 2002EP-0001167
                                                                                                             2001WO-
                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-676582/73
                                                                                                                                                                                                                                                                                              (AEOM-) AEOMICA INC
                                                                                                                                                    30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                30-JAN-2001;
                                                                                        30-JAN-2001:
                                                                                                                                                                                                                                                                                                                                            Zhan J;
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Gaps .,0 0; Indels .0%; Score 13; DB 1; Length 17; 0.0%; Pred. No. 2.5e+02; LDLR cDNA amplifying RT-PCR primer, LDLR/pl. / Match Local Similarity 100.0%; Pred. NO. 2... AAD33183 standard; DNA; 17 BP. 522 CCTGCCGGAGGAG 534 01-JUL-2002 (first entry) AAD33183; Query Match Matches RESULT 464 AAD33183 à g

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Phytanic acid; non-insulin dependent diabetes mellitus; NIDDM; obesity; glucose tolerance; food supplement; feed supplement; hyperinsulinaemia; hypertripidaemia; hypertrension; insulin therapy; hypercholesterolaemia; hypertridyceridaemia; primer; RT-PCR; LDLR; reverse transcription PCR; low-density lipoprotein receptor; ss.

Unidentified

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Sequence 17 BP; 2 A; 6 C; 6 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human IL-2 exon 3-specific probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Page 9; 29pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYMA-) UNIV MARYLAND BALTIMORE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95WO-US04094.
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94US-0219831.
                                                                                                                               30-JUL-2001; 2001EP-0118230.
                                                                                                                                                                                  04-AUG-2000; 2000EP-0116848.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT05895 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           457 GTGGTCAGCAGCC 469
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                                                                                                                                                                                                                                        (ROCH-) ROCHE VITAMINS AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                              Heim M,
                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-270864/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9527052-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAR-1994;
                                                                                                                                                                                                                                                                                              Fluehmann B,
                   EP1177789-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-0CT-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13;
                                                                            06-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    probe; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ~
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This sequence represents an antisense inhibitor of the invention, and inhibits the expression of the human G-alpha-13 protein. The antisense compounds of the invention are of 8 to 30 nucleobases in length, that inhibits the expression of the human G-alpha-13. The antisense compound is useful for treating an animal, particularly humans, having or being prone to a disease or condition associated with the expression of
                                                                                                                                                                                                   Cytokine-specific probes were used to screen products obtd. by RT-PCR amplification (see AATOS878-93) of cellular RNA from human peripheral blood monounclear cells stimulated with an anti-CD3 MAD. Probes were human IL-2 exon 2-specific (AATOS894), human IL-2 exon 3-specific (AATOS895), human IL-5 exon 1/exon 3 junction-specific (AATOS895), human IL-5 exon 1/exon 3 junction-specific (AATOS897) and human GM-CSF exon 1/exon 3 junction-specific (AATOS899) and IL-2 (AATOS899). Novel splice variants of IL-4 (AATOS899) and IL-2 (AATOS900) lacking exon 2 were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  G-alpha-13; human; inhibitor; cancer; antisense compound; therapy; ss.
                                                                            New human interleukin alternative splice variants - expressed by exon(s) 1, 3 and 4 of \rm IL-4 or \rm IL-2, used for regulating the activity of corresp. interleukin(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense compound inhibiting expression of human G-alpha-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human G-alpha-13 antisense inhibitor ISIS# 20763.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Le
2.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 4 A; 4 C; 6 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.0%; Score 13; DB 100.0%; Pred. No. 2.7 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 11; Column 39; 38pp; English.
                                                                                                                                                              Example 6; Page 29; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ31808 standard; DNA; 18 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-JAN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 1.0
Best Local Similarity 100.
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTTCCTCCAGAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-633376/54.
                                       WPI; 1995-358629/46.
Alms W, White B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                              identified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ð
                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to the use of phytanic acid or its derivatives for the treatment or prevention of diabetes mellitus. The invention also relates to a method for treating or preventing non-insulin dependent diabetes mellitus (NIDDM) or other conditions associated with impaired glucose tolerance such as obesity using phytanic acid, their is derivatives or their precursors are useful as pharmaceutical compounds or supplements to foods or feeds for the treatment or prevention of type II or NIDDM, hyperlapidaemia, hypercholesterolaemia, hyperinsulinemia, syndrome X, hypertension, hypertrighyceridaemia, impaired glucose tolerance and related obesity. They are also useful in insulin therapy in combination with known active compounds. The present sequence is low-density continued in hypercesset remascription PCR (RT-PCR) primer used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                            New composition comprising phytanic acid or its derivatives, useful for treating or preventing non-insulin dependent diabetes mellitus, impaired glucose tolerance and related obesity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
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Pred. No. 2.5e+02;
                                                                                                                                                                                                                               Weber P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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100.0%; Pre
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Seguence 18 BP; 4 A; 7 C; 6 G; 1 T; 0 other;

G-alpha-13, such as cancer.

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Gaps

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17 TGCCTTTTTTCT

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                              Human biallelic marker downstream amplification primer SEQ ID NO:8784.
                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N.B. The SEQ ID NOs 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the present invention.
                                                                                                                                                                                                                                                                                                                           Human genome; biallelic marker; high density diseguilibrium map; genomic map; haplotype; polymorphic base; genotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; NPP; PCR primer; amplification; single nucleotide polymorphism; NPP; PCR primer;
                                             ..
          Length 18;
                                         0; Indels
          DB 1; Le 2: 2.7e+02;
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1.0%; Scor.
100.0%; Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 8; Page 2103; 2745pp; English.
                                                                                                                                                                                         AAZ74428 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-IB00822.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-0082614.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-0109732
                                                                           718 GCCCAGCAGCAGG 730
                                                                                                                                                                                                                                                             (first entry)
Query Match
Best Local Similarity 100.0
Matches 13; Conservative
                                                                                                4 GCCCAGCAGCAGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   map of the human genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-013267/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9954500-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                                             10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-OCT-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treatment.
                                                                                                                                                                                                                             AAZ74428;
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                                                                                                                                                                            AAZ74428/C
ID AAZ74
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The present sequence is that of fluorescent target oligonucleotide
T2, which was used to demonstrate an electron hybridisation method
cof the invention. Mn203 stabilised n-type silicon photoelectrodes
coated with a streptavidin-agarose permeation layer were shown to
cligonucleotides by electron hybridisation layer were shown to
cligonucleotides by electron hybridisation. In this process, a
set of unlabelled oligonucleotides (capture strands) are first
targeted to specific locations and anchored. A second set of
fluorescent labeled oligonucleotides (target strands) is then
targeted to the same locations and anchored. A second set of
capture strands. In the example provided, 2 sets of biotinylated
capture probes. Cl (see AAH26908) and C2 (see AAH26909), were
capture probes. Cl (see AAH26908) and C2 (see AAH26909), were
successively transported and anchored to 4 different locations on a
streptavidin-agarose and Mn203 coated amorphous silicon substrate.
Ta (present sequence), were then transported to a location with
complementary capture probes and a location with non-complementary
complementary capture probes and a location of the nation between signal and non-specific background was better
fluorescence signals at the 2 locations with matching sequences.
The ratio between signal and non-specific background was better
capture probes for photoelectrophoretic transport and hybridisation of
manufacture of micro electrophoretic transport and hybridisation of
manufacture of micro electrophoretic transport and hybridisation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Device for photoelectric transport of charged materials in liquid environment for micro- and opto- electronic devices, has a substrate generating light induced current, conductor, permeation layer and light source to illuminate elbetrate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oligonucleotides. The techniques of the invention have wide use ir manufacture of micro electronic and opto electronic devices. Self-assembly fabrication techniques based on DNA polymers enables micron, sub-micron or nanoscale devices to be fabricated.
                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "fluorescent dye with 493 nm absorption and 503 nm emission"
                                                                                                                                                                        Fluorescent oligonucleotide target for electronic hybridisation.
                                                                                                                                                                                                                   Capture probe; hybridisation; electronics; photonics;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           я,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Formosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 60; 119pp; English.
                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                             /*tag= a
/mod_base= "OTHER"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gurtner C,
                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-JAN-2001; 2001WO-US00926.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-JAN-2000; 2000US-0489855
                                         AAH26911 standard; DNA; 18
                                                                                                                            21-DEC-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                         /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Edman CF, Heller MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NANO-) NANOGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-607116/69.
                                                                                                                                                                                                                                        nanotechnology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200153799-A1.
                                                                                                                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-JUL-2001.
                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                     AAH26911;
RESULT 468
                       AAH26911
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New heterocycle derivatives, useful for modulating the activity of RNA
                                                                                                                                                                                                                                      Nucleic acid activity modulator; targetting portion; reactive portion;
                                                       Gaps
                                                      ö
                               Length 18;
                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kawasaki
                              1.0%; Score 13; DB 1; Le
100.0%; Pred. No. 2.7e+02;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Guinosso CJ, Acevedo OL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 5 A; 3 C; 8 G; 1 U; 1 other;
           BP; 5 A; 6 C; 3 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                    /*tag= a
/note= "Optionally absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 136; Column 66; 54pp; English.
                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                   Y strand RNA oligonucleotide #6.
                                                                                                                                                   BP.
                                                                                                                                     92US-0854634.
90US-0566977.
91US-0463358.
                                                                        84 ATAGCAGTTCTAC 96
                                                                                             Ŋ
                                                     Conservative
                                                                                             17 ATAGCAGTICTAC
                                        Local Similarity
Les 13; Conserv
           Sequence 18
                               Query Match
                                                                                                                             RESULT 469
                                                   Matches
X S
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The invention relates to a set of novel map-related biallelic markers, preferably located on obesity disorder-associated chromosomal regions on ehromosomes 3, 10 and 19. The markers are useful for genotyping or estimating the frequency of an allele in a population, for detecting an association between a genotype or haplotype and a phenotype, e.g. a disease involving drug responses, obesity or disorders related to obesity, such as hyperuricaemia, digestive pathology, hepatic function disorders, cancer, cardiovascular disease, hypertension, hyperlipidaemia, insulin disorders, atheromatous disease and cardiac insufficiency. The markers are useful for detecting a statistical correlation between a biallelic marker allele and a phenotype and/or between a biallelic marker haplotype and a phenotype and/or between a biallelic marker haplotype and a phenotype and/or between a biallelic marker haplotype and a phenotype sente represents a PCR primer used to amplify a human obesity-associated biallelic marker.
                                                                                                                                                                                                                                           Human; obesity associated-biallelic marker; chromosome 10; obesity; ss; delug response; hyperuticaemia; digestive pathology; hypertension; cancer; hepatic function disorder; cardiovascular disease; hyperlipidaemia; PCR; insulin disorder; atheromatous disease; cardiac insufficiency; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Set of novel map-related biallelic markers, preferably located on obesity disorder-associated chromosomal regions on chromosomes 3, 10 and 19, useful, for e.g. detecting statistical correlations between marker allele and a phenotype -
                                                                                                                                                                                                         Human obesity-associated biallelic marker downstream PCR primer #57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bihain B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chumakov I, Abderrahim H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Pred. No. 2.7e+02; ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .0%; Score 13; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 6 A; 3 C; 5 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 274; 311pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                           BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-JUN-2001; 2001WO-IB01477.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-JUL-2000; 2000US-219704P.
                                                                                           ABK41151 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ68252 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        325 CTGCATCATCCTG 337
                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTGCATCATCCTG 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cohen D, Blumenfeld M,
CCTGGCCTTNCCAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003 (updated)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-155043/20.
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Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                WO200206525-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                 21-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                      24-JAN-2002
                                                                                                                                 ABK41151;
18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 471
AAQ68252/C
ID AAQ682
XX
AC AAQ682
XX
DT 25-MAR
                                                       RESULT 470
                                                                           ABK41151/c
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1.0%; Score 13; DB 1; Length 18; 92.9%; Pred. No. 2.7e+02; tive 0; Mismatches 1; Indels

Query Match
Best Local Similarity 92.9
Matches 13, Conservative

1057 CCTGGCCTTCCCAT 1070

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The method aims to provide a collection of highly reproducible microsatellite marker sequences (MMS) at approx. 10-50 cM intervals throughout the human genome which can be detectably labelled. The MMS are polymorphic, simple sequence repeats and can be used in automated genotyping. esp. fluorescence-based. The primers correspond to the unique DNA sequence surrounding each marker, and PCR is used to
                                                                                                                                                   Five sets of purine-rich methylphosphonate oligonucleosides ("MP oligomers") and complementary ribooligonucleosides ("RNA oligomers") were examined for their ability to form triple helix complexes. (Set 1:G2100 and R39; Set 2:G2101 and R289; Set 3:G2102 and R291; Set 4:G2104 and R293; Set 5:G2106 and R84 - see AAG68229-34 and AAQ80860-63).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kit for automated genotyping contg. pairs of PCR primers - designed to amplify polymorphic nucleotide repeat sequences, arranged in sets each with a characteristic fluorescence label, useful e.g. in detection of disease related genetic rearrangement
                                                                                                                                                                                                                                                                           It was shown that replacing two adenines with two thymidines is somewhat destabilising, and that replacing two guanines with two cytidines is more destabilising w.r.t. triple helix formation (Updated on 25-MAR-2003 to correct PN field.)
                                 Detection, recognition, inhibition and alteration of single and double stranded target nucleic acid sequences - by formation of striple helix structure using 2 oligomers which block translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer; polymerase chain reaction; PCR; linkage study; locus;
microsatellite marker sequence; automated genotyping; allele;
polymorphism; detection; Homo sapiens; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                      0.9%; Score 12.8; DB 1; Length 16;
87.5%; Pred. No. 2.5e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer A (Group 11, set B) for marker D5S425, chromosome
                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16 BP; 8 A; 2 C; 6 G; 0 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure, Fig 7K-2, 104pp, English.
                                                                                                                   Example 4; Page 39; 67pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1216 TTCCCTGTACATTTGT 1231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94WO-US13945.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ95859 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16 rrcccrercccrrrer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-FEB-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1995-215278/28.
WPI; 1994-217542/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9515400-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-JUN-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Levitt RC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ95859;
                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ95859/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ΩD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Triple helix formation with 2:1 MP:RNA oligomers was demonstrated with thermal denaturation methods. Exemplary triple helix forming MP-oligomers are given in AAQ66242-52. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detection, recognition, inhibition and alteration of single and double stranded target nucleic acid sequences - by formation of a triple helix structure using 2 oligomers which block translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Purine, methylphosphonate, MP; triple helix, translation, oligonucleoside; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                          Triple helix forming methylphosphonate oligomer 2104.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Purine-rich methylphosphonate oligonucleoside G2104.
                                                                                Methylphosphonate; MP; triple helix; translation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.9%; Score 12.8; DB 1; Best Local Similarity 87.5%; Pred. No. 2.5e+02; Matches 14; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 16 BP; 8 A; 2 C; 6 G; 0 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 11; Page 50; 67pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93WO-US11986
                                                                                                                                                                                                                                                                                                       92US-0987746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16 rrecererecerrrer 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1216 TICCCIGIACATITGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reynolds MA;
                                                                                                                                                                                                                                                                                                                                                                                     Reynolds MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (updated)
    (first
                                                                                                       oligonucleoside; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1994-217542/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GENT-) GENTA INC.
                                                                                                                                                                                                                                                                                                                                             (GENT-) GENTA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9413326-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-DEC-1993;
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                                                                                                                                                                                                                                                                                                       08-DEC-1992;
                                                                                                                                                                                   WO9413326-A1
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16-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arnold LJ,
                                                                                                                                                                                                                         23-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                     Arnold LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ80862;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ80862/
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(ie. a difference in the number of repeats) between individuals, the markers can be particularly informative. The MMS can be ideal for linkage studies. Kits comprise at least 4 groups, of at least 3 sets, each comprising labelled primers for PCR amplification of the DNA. Group 11 primer pairs are shown in AAQ5881-82. The published size range of the D55425 allele is 224-248 bp, and the degree of heterozygosity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the following microsatellite markers on chromosome 1q21q31 associated with occurrence of juvenile glaucoma: afm350yhl; afm122xa3; nga1; afm218xb3; afm218ye5; afm218xb10; afm157xe7 and NGA5. An oligonucleotide primer of the present sequence was used with a primer having the sequence given in AAT43026 to amplify the afm350yhl marker. Apart from detecting predisposition to disease, the microsatellites should allow localisation, and thus isolation, of
 each polymorphism. When the MMS show considerable polymorphism
                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detecting pre-disposition to juvenile glaucoma - from presence of specific micro:satellite markers on chromosome 1q21q31, also DNA from the region defined by these markers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Predisposition to juvenile glaucoma is detected by characterising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Microsatellite, genetic marker, screening, detection, PCR primer, polymerase chain reaction, juvenile glaucoma, predisposition, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Juvenile glaucoma marker afm350yhl upstream amplification primer.
                                                                                                                                                                                                                      ..
                                                                                                                                                                                    0.9%; Score 12.8; DB 1; Length 16; B7.5%; Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= a
/note= "linked to JOE fluorochrome label"
                                                                                                                                                                                                                      2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (INRM ) INSERM INST NAT SANTE & RECH MEDICALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 16 BP; 2 A; 9 C; 1 G; 4 T; 0 other;
                                                                                                                                                  Sequence 16 BP; 2 A; 8 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the gene involved in juvenile glaucoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                 in the population is about 77%.
                                                                                                                                                                                                                                                                                                                                                                        ВЪ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example; Page 7; 25pp; French
                                                                                                                                                                                                                                                     459 GGTCAGCAGCCTGCAG 474
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96WO-FR00592.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95FR-0004590.
                                                                                                                                                                                                                                                                                    GGTCAGCAGGCTGGAG 1
                                                                                                                                                                                                                                                                                                                                                                       AAT43025 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Garchon HJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1996-485791/48.
                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9633287-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                        19-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-OCT-1996,
                                                                                                                                                                                                                      14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bach JF,
                                                                                                                                                                                                                                                                                     16
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                                                                                                                                                                                                                      Matches
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0.9%; Score 12.8; DB 1; Length 16;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This sequence represents a primer of the invention. The invention relates to sequences of at least two mucleotides of formula:

(X)m5'-(alpha)n-beta-N3'; or (X)m5'-(gamma)k-delta-N3'; where
X = a labelled compound and/or a nucleotide with voluntary sequence;
m = 0 or 1; alpha = thymine; n = natural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, guamine or cytosine;
N = adenine, gamma = thymine;
k = natural number of 3 or over indicating the repetition of gamma, in which thymine expressed by gamma is composed of 1/3 or less of adenine, guanine and/or cytosine. The new nucleotides are useful as primers for RT-PCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene sequences.
                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                             RT-PCR primer; DNA sequence determination; gene sequence analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             - useful as primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              o.
                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide #31 used in a method for primer selection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                Indels
Pred. No. 2.5e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 12.8; DB 1;
17.5%; Pred. No. 2.5e+02;
.ve 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 16 BP; 1 A; 0 C; 1 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptides having at least two new nucleotides
                                                                                                                                                                                                                                 RT-PCR primer of the invention SEQ ID 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 10; 19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1144 TTTTTTTTTTGGA 1159
                                          1042 TCTTCCCACGACAGCC 1057
                                                                                                                                             AAX18366 standard; DNA; 16 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16
 87.5%;
                                                                                                                                                                                                                                                                                                                                                                             97JP-0208312.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAC63258 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                    (first entry)
                                                                   1 TCTTCCCACCACTGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 87.5
Matches 14; Conservative
                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                    (TAKI ) TAKARA SHUZO CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-183822/16.
   Best Local Similarity
                                                                                                                                                                                                                                                                                                                    JP11032765-A
                                                                                                                                                                                                                                                                                                                                                                             18-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                         18-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-FEB-2001
                                                                                                                                                                                                    11-MAY-1999
                14;
                                                                                                                                                                                                                                                                                                                                                09-FEB-1999
                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                        AAX18366;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAC63258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RT-PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 476
                                                                                                                RESULT 475
                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC63258,
                                                                                                                              AAX18366
                                                                                                                                                          g
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Thu Jan

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The present invention relates to a method for selecting PCR primers for nucleic acid amplification. The method comprises determining the melting temperature (T m) range for degenerate oligonucleotide primers with a fixed-sequence portion (FS) and a degenerate sequence portion (DS) by searching known portion of a nucleic acid template for a sequence complementary to a desired FS of a primer. Nucleotide base pairs flanking or interspersed between the sequence complementary to a DS of one of the primers are detected and T m is calculated. The method of the present invention allows primers which produce more efficient DNA amplification to be produced. The present sequence is a primer used in the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The specification describes methods for genotyping polymorphisms that result in the gain or loss of an endonuclease site polymorphisms (ESPS). The method comprises deriving a set of concomitantly amplifiable target DNA fragments from sample DNA, treating the target DNA fragments with a
                                                        Determining Tm range for several degenerate primers with a fixed-sequence and a degenerate-sequence portion for use in polymerase chain reaction amplification by identifying a specific sequence in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel method for restricted amplicon analysis of endonuclease site polymorphisms resulting in loss of restriction sites, useful for multiplex genotyping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphism; endonuclease site polymorphism; ESP; genetic marker; restriction endonuclease; high throughput genetic analysis; animal breeding; plant breeding; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.9%; Score 12.8; DB 1; Length 1
87.5%; Pred. No. 2.5e+02;
ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer used for screening for ESP carrying fragments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 16 BP; 1 A; 9 C; 6 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page 34; 67pp; English.
                                                                                                                                                                        Disclosure, Fig 3B; 34pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCGCCCACCCGGACGC 206
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA46382 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-SEP-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cceccceccceaegc
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Т
                                                                                                                                nucleic acid template
                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-376586/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (METH-) METHEXIS NV.
                WPI; 2000-656235/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
ses 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200028081-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-MAY-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zabeau M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA46382;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a method for selecting PCR primers for nucleic acid amplification. The method comprises determining the melting temperature (T_m) range for degenerate oligonucleotide primers with a fixed sequence portion (FS) and a degenerate-sequence portion (DS) by searching known portion of a nucleic acid template for a sequence complementary to a desired FS of a primer. Nucleotide base pairs flanking or interspersed between the sequence complementary to a DS of one of primers are detected and T m is calculated. The method of the present thouse primers which produce more efficient DNA amplification to be produced. The present sequence is a primer used in the method of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                Determining Tm range for several degenerate primers with a fixed-sequence and a degenerate-sequence portion for use in polymerase chain reaction amplification by identifying a specific sequence in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer; nucleic acid amplification; melting temperature; T_m; 88.
PCR primer; nucleic acid amplification; melting temperature; T_m; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
0.9%; Score 12.8; DB 1; Length 16;
Best Local Similarity 87.5%; Pred. No. 2.5e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide #73 used in a method for primer selection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16 BP; 0 A; 4 C; 11 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 3B; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TECHNOLOGIES LLC
                                                                                                                                                                                                                                                                      TECHNOLOGIES LLC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    191 CCGCCCACCCGGACGC 206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-0127891.
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                                                                                                                                                                                                                          99US-0127891
                                                                                                                                                                           05-APR-2000; 2000WO-US08962
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acid template
                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-656235/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENO-) GENOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200060123-A2
                                                                                      WO200060123-A2
                                                                                                                                                                                                                                                                    GENOME
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                                              Homo sapiens.
                                                                                                                                                                                                                          06-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-OCT-2000
                                                                                                                                                                                                                                                                                                                Senapathy P;
                                                                                                                                12-OCT-2000
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0;

Gaps

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probe restriction endonuclease reagent, amplifying the DNA fragments, and analysing the DNA to determine which target fragments have been, and have not been, amplified. Target DNA fragments which have been amplified lack a recognition site for the probe restriction endonuclease reagent, and target fragments having a recognition site for the probe restriction disagnosing the immense number of genetic markers that are needed to unravel complex traits. The method is capable of unravel complex traits. The method is useful for high throughput genetic identify genes involved in quantitative agronomic traits. The methods are also useful to monitor mutations in specific genes or loci in addition to scanning the entire genome. The present sequence represents a for a per primer used in the course of the invention, for genetic analysis
         $$$$$$$$$$$$$$$$$$$$$$$$$$$
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BP; 3 A; 4 C; 5 G; 4 T; 0 other; Sequence 16

Length 16; Indels Score 12.8; DB 1; L Pred. No. 2.5e+02; ^. Mismatches 2; 0; Mismatches 739 CTGCCGCATGTTGCTG 754 0.9%; 76 Local Similarity 87. Query Match Best Loca Matches ò

CTGACGCATGTTGCAG

RESULT 479

AAF73460 standard; DNA; 16 BP AAF73460

AAF73460;

(first entry) 08-MAY-2001

HGF nucleic acid ligand SEQ ID NO: 10.

Hepatocyte growth factor/ scatter factor; HGF; c-met; integrin; stroke; cell adhesion; cell migration; nucleic acid ligand; thrombosis; cancer; hypertension; arteriosclerosis; myocardial infarction; restenosis; rheumatoid arthritis; macular degeneration; endometriosis; psoriasis; osteoporosis; DNA-RNA hybrid; ss.

Synthetic

/note= "all bases are 2'OMe" Location/Qualifiers "OTHER" 5..16 /\*tag= a 1..16 /\*tag= b /mod\_base= modified base misc\_RNA

WO200109159-A1.

08-FEB-2001

24-JUL-2000; 2000WO-US20139.

99US-0364539. 99US-0364543. 29-JUL-1999; 29-JUL-1999;

(NEXS-) NEXSTAR PHARM INC.

Lochrie M; Stephens A, Janjic N, Rabin R, Gold L, ٠, Ruckman

WPI; 2001-103180/11.

Isolation of nucleic acid ligands to hepatocyte growth factor, its receptor c-met and integrins, useful for treating tumors, deep vein thrombosis and diabetic retinopathy -

Example 1; Fig 2; 226pp; English

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present invention provides nucleic acid ligands to hepatocyte growth or/scatter factor (HGF), its receptor c-met and integrins. Integrins
                      factor/scatter factor (HGF), its receptor c-met and integrins. Integrins are involved in cell adhesion and migration, and HGF is a cytokine involved in cell proliferation and migration. The ligands of the invention are useful in the treatment of diseases such as cancer, thrombosis, hypertension, arteriosclerosis, myocardial infarction, rheumatoid arthritis, macular degeneration, endometriosis, psoriasis, stroke, osteoporosis and restenosis. The present sequence is an example
                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                ó
                                                                                                                                                                                                                                                                                                Length 16;
                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                               Sequence 16 BP; 1 A; 4 C; 7 G; 2 T; 2 U; 0 other;
                                                                                                                                                                                                                                                                                            0.9%; Score 12.8; DB 1;
75.0%; Pred. No. 2.5e+02;
cive 2; Mismatches 2;
                                                                                                                                                                                               of a ligand of the invention.
                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                           12;
                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                        Matches
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297 GTCTGCTGTGGGGCT 312 16 1 GrérécudAscesser ò g

; 0

Gaps

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ABL95939 standard; DNA; 16

ABL95939;

(first entry) 19-JUN-2002 Probe #23 for assaying nucleic acids.

Probe; polymorphism detection; mutation detection; disease diagnosis; microbial identification; ss.

Unidentified.

WO200208414-A1.

31-JAN-2002.

27-JUN-2001; 2001WO-IB01147.

27-JUN-2000; 2000JP-0193133. 03-AUG-2000; 2000JP-0236115. 26-SEP-2000; 2000JP-0292483.

IT ADVANCED IND SCI & TECHNOLOGY. INST (KANK-) KANKYO

Yamada K; Kurata S, Torimura M, Kanagawa T, Kamagata Y, Kurane R, Yokomaku T;

WPI; 2002-195876/25.

Fluorescently-labeled nucleic acid probes for assaying nucleic acids and their polymorphism and mutation, particularly useful in science and medicine for e.g. analytical applications, disease diagnosis and microbial identification

Example 22; Page 76; 152pp; Japanese.

The present invention relates to nucleic acid probes, which are useful for assaying nucleic acids by hybridising with a target nucleic acid, in which a single-stranded oligomuchecide is labelled with a fluorescent substance and a quencher in a manner that the fluorescence intensity of the hybridisation reaction system is increased after completion of the hybridisation but no stem loop structure is formed. The probes are useful for assaying nucleic acids and their polymorphism and mutation, particularly useful for e.g. analytical applications, disease diagnosis and microbial identification. The present sequence was used to illustrate the invention. 

Sequence 16 BP; 0 A; 6 C; 2 G; 8 T; 0 other;

us09904568-3.rng

Diagnostic; Gram-positive bacterium; high G+C content; amplification; mycobacterial infection; PCR; primer; probe; detection; ss.

Corynebacterium jeikeium

WO200297126-A2

05-DEC-2002

23S rDNA helix 54 region probe SEQ ID 33.

(first entry)

10-JUN-2003

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                                                                                                                                                                                                                                                                                                             Bronchial asthma attack; SCCAI; SCCA2; gene expression; risk of attack;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to a method of examining attacks of bronchial asthma by using SCCA1 and/or SCCA2 as the indicator genes. The method comprises determining the expression levels of the indicator genes in the biological sample from a patient, and comparing the expression level with that in the sample of a healthy individual. The method is useful for examining bronchial asthma, which is useful in assessing the risk of attacks with SCCA1 and SCCA2 gene expression levels as indication. The method is cheap and capable of operating in high throughput, even for bedside diagnosis. The present sequence represents a PCR primer used in the examples of the present invention.
                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Izuhara K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Method for examining bronchial asthma based on SCCA1 and SCCA2 as allergy-associated genes, useful in assessing risk of attacks with their expression levels as indication
                                       .;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%; Score 12.8; DB 1; Length 16; 87.5%; Pred. No. 2.5e+02; Live 0; Mismatches 2; Indels
     Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matsui K, Yoshida N, Sugita Y, Hamasaki Y,
                                    2; Indels
0.9%; Score 12.8; DB 1;
ilarity 87.5%; Pred. No. 2.5e+02;
Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 16 BP; 3 A; 6 C; 4 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; Page 20; 44pp; Japanese.
                                                                                                                                                                                                                                                                              Human SCCA2 gene, PCR primer #2.
                                                                 1214 CCTTCCCTGTACATTT 1229
                                                                                           1 CCTTCCCTGTTT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-AUG-2002; 2002WO-JP07918.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-AUG-2001; 2001JP-0239857
                                                                                                                                                                               ABX94194 standard; DNA; 16
                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GENO-) GENOX RES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-248304/24.
                 Best_Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                              WO2003014395-A1
                                                                                                                                                                                                                                                                                                                              PCR; primer; ss
                                                                                                                                                                                                                                             10-JUN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ohtani N,
                                                                                                                                                                                                                ABX94194;
   Query Match
                                                                                                                                              RESULT 481
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                                                                                                                                                                 ABX94194
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Detecting and identifying Gram-positive bacteria of high G/C content, useful particularly for diagnosis of mycobacterial infection, by specific amplification and hybridization

(HAIN-) HAIN LIFESCIENCE GMBH

WPI; 2003-140491/13.

Weizenegger M;

09-APR-2002; 2002WO-EP03956. 03-MAY-2001; 2001DE-1021505. Claim 1b; Figure 2B; 34pp; German.

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This invention describes a novel method for the diagnostic detection and/or identification of Gram-positive bacteria that have a high G+C content, especially Mycobacteria. The method comprises subjecting a sample to nucleic acid amplification using the PCR primers represented in ABX94483-ABX94492. The amplification mixture, or part of it, is then tested for hybridisation to at least one of the probes represented in ABX94493-ABX94524 which can be immobilised on a solid phase or used in kit form. The specified primers/probes provide highly specific detection of particular Gram positive bacteria, which are difficult to differentiate by morphological or biochemical tests and/or those which take a long time to test because of their slow growth.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; ohronic myelogenous leukaemia; CML; cancer; philadelphia chromosome; inflammation; autoimmune disease; atheroselerosis; myocardial infraction; stroke; restenosis; transplant rejection; rheumatoid arthritis; psoriasis; myocardial ischaemia; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rat ICAM hammerhead ribozyme target sequence (nt. position 1006).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 16 BP; 4 A; 6 C; 4 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              598 ACCAGCCTGAAGCCTG 613
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(first entry)
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Best Local Similarity 87.57
Matches 14; Conservative
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AAT53533/c
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828 GCAGCTGAAGCTTTCA 1 GCAGCTGCAGCTTCCA

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ABX94515 standard; DNA; 16 BP

RESULT 482

ABX94515

ABX94515;

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17 CCTGGTGATACTCCCA 2

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AAT53757/c
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RNA at the nucleotide base position indicated in the DE line.

Regions of the mRNA that do not form secondary folding.

Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin structures and that contain potential hammerhead and hairpin structures and that contain potential hammerhead and hairpin structures and that contain severe identified by computer analysis.

Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in patients with renumatoid arthritis, asthma and other inflammatory disorders.

(Updated on 25-MAR-2003 to correct PI field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ribozymes having modified bases and methods for producing them for use in inhibiting disease related genes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Page 202; 407pp; English.
                                                                                                                                                                                                                                                                  95US-0380734.
94US-0201109.
94US-021109.
94US-02227954.
94US-0228041.
94US-024804.
94US-0291932.
94US-0291433.
94US-0291433.
94US-0291433.
94US-0291433.
94US-0291433.
94US-0291433.
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94US-0316771.
94US-0319492.
94US-0321993.
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94US-0337608.
94US-0345516.
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                                                     Rattus rattus,
                                                                                                        WO9523225-A2
                                                                                                                                                                                                                23-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                   15-APR-1994;
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23-SEP-1994;
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AIDS; ss.
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Bnzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; ref A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bor-abl; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; philadelphia chromosome; inflammation; autoimmune disease; atherosclerosis; myocacdial inflamction; stroke; restenosis; transplant rejection; rheumatoid arthritis; psoriasis; myocardial ischaemia; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stinchcomb DT, Chowrira B, Direnzo A, Draper KG, Dudycz LW; Grimm S, Karpeisky A, Kisich K, Matulic-adamic J, Mcswiggen JA; Modak A, Pavco P, Beigleman L, Sullivan SM, Sweedler D; Thompson JD, Tracz D, Usman N, Wincott FE, Woolf T;
                                                                                      Rat ICAM hammerhead ribozyme target sequence (nt. position 2607).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ribozymes having modified bases and methods for producing them for use in inhibiting disease related genes
AAT53757 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                    95US-0380734.
94US-0211109.
94US-0221934.
94US-0222483.
94US-0227958.
94US-0227958.
94US-0221932.
94US-0291433.
94US-02914397.
94US-0311749.
94US-0311749.
94US-0311749.
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                                                               (first entry)
                                                   (updated)
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                                                                                                                                                                                                                                                                    Rattus rattus.
                                                  25-MAR-2003
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                                                                                                                                                                                                                                                                                                                                            23-FEB-1995;
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06-JUL-1994;
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                          AAT53757;
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Gaps

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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (Preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour anglogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
    smooth muscle cell hyperproliferation in restenosis, especially after coronary angioplasty, and in cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vascular endothelial growth factor receptor; VBGF receptor; flt-1; flk, hammerhead ribozyme; halripin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; coular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene expression or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse flt-1 VEGF receptor hammerhead ribozyme substrate #702.
                                                                                                                         Length 17;
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                                                                                           Score 12.8; DB 1; Le
; Pred. No. 2.7e+02;
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0.9%; Score 12.8; DB 1;
Best Local Similarity 56.2%; Pred. No. 2.7e+02;
Matches 9; Conservative 5; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stinchcomb
                                                                       Sequence 17 BP; 1 A; 8 C; 3 G; 5 U; 0 other;
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                                                                                                                                                                        4; Mismatches
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                                                                                                                         0.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-JUL-1999 (first entry)
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                                                                                                                                                                      10; Conservative
                           coronary angioplasty,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-259017/23.
                                                                                                                         Query Match
Best Local Similarity
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                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human c-myb hammerhead ribozyme target sequence (nt. position 1276).
                                     The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves ICAM-1 mRNA at the nuclectide base position indicated in the DE line. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes cleave the ICAM-1 target sequences and thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in patients with theumatoid arthritis, asthma and other inflammatory disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New enzymatic nucleic acid molecules - which cleave RNA produced by e.g. c-myb, for treating restenosis or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ÷
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bnzymatic nucleic acid; hammerhead; ribozyme; cleavage; human;
smooth muscle cell; hyperproliferation; restenosis; cancer;
c-myb; coronary angioplasty; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.9%; Score 12.8; DB 1; Best Local Similarity 87.5%; Pred. No. 2.7e+02; Matchés 14; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 4 A; 3 C; 7 G; 3 U; 0 other;
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Claim 2; Page 204; 407pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 68; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          334 CCTGGTGATAGTCACA 349
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94US-0245466.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JAN-1995;
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Gaps

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Zea mays.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Loca
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (WGF). A patient (preferably human) having a condition associated with the level of the fins-like tyrosine kinse 1 (fit-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (fik-1) (e.g. tumour anglogenesis, ocular diseases, psoriasis and rhumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                   vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour anglogenesis; psoriasis; hetumatoid archritis; coular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
                                                                                                                                                                                                                                                                                                                                                                       mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                          Human KDR VEGF receptor hammerhead ribozyme substrate #636.
                                                                                                                                                                                                                                                                                                                                  Stinchcomb D;
                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid molecule modulating VEGF receptor(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 2 A; 3 C; 5 G; 7 U; 0 other;
                                                                                                                                                                                                                                                                                                                                 McSwiggen J, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; Page 116; 218pp; English.
                                                AAX71624 standard; RNA; 17 BP
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CUUCGCAUACUGCUCA 16
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                                                                                                                                                                                                                                                                                                   CHIRON CORP.
RIBOZYME PHARM INC.
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                                                                                       (first entry)
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les 14; Conservative
                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-259017/23.
                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                Escobedo J,
                                                                                                                                                                                                          WO9715662-A2
                                                                                                                                                                                                                                                    25-OCT-1996;
                                                                                                                                                                                                                                                                     11-JAN-1996;
26-OCT-1995;
                                                                                       28-JUL-1999
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                                                                                                                                                                                                                                01-MAY-1997
                                                                   AAX71624;
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(RIBO-)
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                                     AAX63011
ID AAX6
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AC AAX6
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DT 16-J
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The present invention describes an enzymatic nucleic acid molecule (I) with RNR cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDNA sequence encoding maize Delta-9 described is a gene comprising a cDNA sequence encoding maize preferably Delta-9 desaturase or a granule bound starch synthase (GBSS) gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate caffeine synthesis in a coffee plant, nicotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, chrysanthemum or marigold plant or lignin production in a tobacco, petunia, aspen, poplar or pine plant.
                                                             Maize; corn; Zea mays; delta-9 desaturase; GBSS; target; substrate; granule bound starch synthase; hammerhead ribozyme; hairpin ribozyme; modulation; gene expression; transgenic plant; cleavage; canola plant; caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Maize; corn; Zea mays; delta-9 desaturase; GBSS; target; substrate; granule bound starch synthase; hammerhead ribozyme; hairpin ribozyme; modulation; gene expression; transgenic plant; cleavage; canola plant; caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound starch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Granule bound starch synthase hammerhead substrate SEQ ID NO:75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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Delta-9 desaturase hamerhead ribozyme target SEQ ID NO:886.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Merlo DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   McSwiggen JA,
Zwick MG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 2 A; 1 C; 2 G; 12 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 38; Page 87; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   O, Guo L,
Young SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1103 ATTATGTAGTTTTCTG 1118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              synthase in maize or canola
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(RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Skokut TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-202224/18.
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                                                                                                                                                                                                                                                                                                                                                 WO9710328-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-JUL-1996;
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Merlo PAO, S
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The present invention describes enzymatic nucleic acid molecules (NAMs) which specifically cleave RNA derived from an epidermal growth factor receptor (EGF-R) gene. AAV9721 to AAV98043 and AAV9899 to AAV99090 represent specifically claimed target sequence from human EGF-R. AAV98044 to AAV98866 and AAV98867 to V9878 represent hammerhead ribozymes and haippin ribozymes respectively for human EGF-R. The NAMs are useful for cleaving EGF-R RNA in the treatment of a condition associated with EGFR expression levels e.g. to inhibit cell proliferation in the prevention or texament of cancers. The NAMs can also be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of EGF-R RNA in a cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence invention describes ribozymes targeted to modulate the synthesis and/or expression of interleukin (IL)-2R gamma encoded RNA. AAV93889 to AAV94574 represent specifically claimed ribozymes, and AAV94575 to AAV95260 represent specifically claimed substrate sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, IL-2 receptor g-chain, interleukin 2 receptor gamma chain; hammerhead ribozyme; hairpin ribozyme; substrate; expression; cancer; autoimmune disease; psoriasis; allergy; inflammatory disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                   Enzymatic nucleic acids - which cleave RNA derived from an epidermal growth factor receptor, useful for inhibiting cell proliferation and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           - useful for treating e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse IL-2 receptor g-chain substrate position 403.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12.8; DB 1;
Pred. No. 2.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 3 A; 8 C; 3 G; 3 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ribozymes targeted to interleukin 2 - use cancer, autoimmune disease and allergies
      Fell P, McSwiggen JA;
                                                                                                                                                                Claim 5; Page 77; 109pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; Page 41; 61pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-FEB-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               574 CAGCAGGCCCTCCGTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12; Conservative
                                                                                                                          for treating cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             graft rejection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-333332/29.
                                         WPI; 1998-437449/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         McSwiggen JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9824913-A2
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        Akhtar S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAV94931;
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        # X & X # # # X & X O O O O O O O O O O X O
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                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes an enzymatic nucleic acid molecule (I) with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDNA sequence encoding maize Delta-9 desaturase. (I) can be used to modulate expression of a gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate caffeine synthesis in a coffee plant, nicotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a rose, petunia, aspen, poplar or pine plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; epidermal growth factor receptor; EGFR; EGF-R; target sequence; hammerhead ribozyme; hairpin ribozyme; inhibition; cell proliferation; cancer; genetic drift; detection; mutation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                           starch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound synthase in maize or canola
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 17;
                                                                                                                                                                                                                        Merlo DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human EGF-R target sequence nucleotide position 3888.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.9%; Score 12.8; DB 1; 168.8%; Pred. No. 2.7e+02; tive 3; Mismatches 2;
                                                                                                                                                                                                                      McSwiggen JA,
Zwick MG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 1 A; 6 C; 6 G; 4 U; 0 other;
                                                                                                                                                                                                                Folkerts O, Guo L, cokut TA, Young SA,
                                                                                                                                                                                                                                                                                                                                                                                             Claim 41; Page 73; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              891 GCTGCGGTACAGCGTG 906
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                                                                                                                                                   (DOWC ) DOWELANCO.
(RIBO-) RIBOZYME PHARM INC.
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Best Local Similarity 68.8
Matches 11, Conservative
                                                                                                                                                                                                            1 BE, Folker
AO, Skokut TA,
                                                                                                                                                                                                                                                                            WPI; 1997-202224/18.
WO9710328-A2
                                                                        12-JUL-1996;
                                                                                                                  13-JUL-1995;
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                                                                                                                                                                                                              Edington BE,
Merlo PAO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-MAR-1999
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                                   20-MAR-1997
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Humanised anti-HM1.24 antibody PCR primer SEQ ID NO:72.

21-SEP-1998

AAV39410;

BP.

AAV39410 ID AAV39410 standard; DNA; 17

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from the present invention. The ribozymes can be used for the treatment of, e.g. graft rejection, autoimmune disease, cancer, psoriasis, allergy and other inflammatory conditions. The ribozymes are also used to induce tolerance in a recipient to alloantigen from a donor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence invention describes ribozymes targeted to modulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the synthesis and/or expression of interleukin (IL) 2R gamma encoded RNA. AAV93889 to AAV94544 represent specifically claimed ribozymes, and AAV94575 to AAV95260 represent specifically claimed substrate sequences from the present invention. The ribozymes can be used for the treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  allergy and other inflammatory conditions. The ribozymes are also used to induce tolerance in a recipient to alloantigen from a donor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, IL-2 receptor g-chain, interleukin 2 receptor gamma chain,
hammerhead ribozyme; hairpin ribozyme; substrate; expression; cancer,
autoimmune disease; psoriasis; allergy; inflammatory disease;
                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of, e.g. graft rejection, autoimmune disease, cancer, psoriasis, allergy and other inflammatory conditions. The ribozymes are als
                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ribozymes targeted to interleukin 2 - useful for treating e.g. cancer, autoimmune disease and allergies
                                                                                                                      0.9%; Score 12.8; DB 1; Length 17;
81.2%; Pred. No. 2.7e+02;
iive 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%; Score 12.8; DB 1; Length 17; ilarity 75.0%; Pred. No. 2.7e+02; Conservative 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse IL-2 receptor g-chain substrate position 248.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 4 A; 7 C; 4 G; 2 U; 0 other;
                                                                                     Sequence 17 BP; 3 A; 8 C; 4 G; 2 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 41; 61pp; English
                                                                                                                                                                                            625 GACCAGCTCCAGGAGC 640
                                                                                                                                                                                                                                                                                                                   AAV94898 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Stinchcomb DT;
                                                                                                                                                                                                                 GUCCAGCUCCAGGACC 16
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                                                                                                                                                           13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   graft rejection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  McSwiggen JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9824913-A2
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                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                RESULT 492
                                                                                                                                                           Matches
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humanised anti-HM1.24 antibody has been developed which comprises human L and H chain C regions, and L and/or H chain V regions containing material originating in mouse anti-HM1.24 antibody. The V containing material originating in mouse anti-HM1.24 antibody. The V ceptons contain framework (FR) regions of human origin and complimentarity determining regions (CDR) of mouse origin, leading to a reshaped humaning regions (CDR) of mouse origin, leading to a reshaped humaning regions (The C regions are human C (L-chain) and human (especially C gamma 1) (H-chain). The FR regions of the L chain V region are derived from human subtype HSG1 (e.g. from human antibody RE1) and the FR regions of the H chain V region are derived from human antibody HG3 (e.g. FR1-3 from human antibody HG3 (e.g. FR1-3 from human antibody HG3 and FR4 from human antibody JH6). The present sequence represents a PGR primer used in an example from the present invention. The antibodies are used for the treatment of myeloma, especially by injection, intravenously, intravenously
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                                                                                                                                                                                     Mouse; human; humanised; anti-HM1.24 antibody; myeloma; FR; CDR; framework region; complimentarity determining region; antigenicity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            therefore effective therapeutically in humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Integrin alpha 6 subunit substrate sequence SEQ ID NO:3717.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Humanised anti-HM1.24 antibody - for treatment of myeloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tsuchiya M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12.8; DB 1;
Pred. No. 2.7e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 5 A; 7 C; 4 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ono K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 9; Page 140; 210pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ohtomo T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 870 CCCCACAGCCAAGTTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEIYAKU KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kosaka M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-286421/25.
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                                                                                                                                                                                                                                    PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Koishihara Y,
                                                                                                                                                                                                                                                                                                    Mus sp.
Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-OCT-1997;
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                                                                                                                                                                                                                                                                                                                                                                            WO9814580-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yoshimura Y;
                                                                                                                                                                                                                                                                                                                                                                                                                           09-APR-1998,
                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA20491;
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Gaps

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227 CTCAGCCTCAGGCATC 242

12;

Matches

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RESULT 493

us09904568-3.rng

(first entry)

16-JUL-1999

AAX59454;

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Human, aryl hydrocarbon nuclear transport, ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammernead ribozyme; andiogenic factor; cytostatic; antidabetic; ophthalmologic; antiniflammatory; antiarthritic; antipsoriatic; ARWD; dermatological; RNA cleavage, cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; veruca vulgaris; angiofibroma; tuberous solerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss.
                                                                                                                                                                                                                                                                                                    Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors
                                                                                                                                                                                                                                                                                                                                     Claim 55; Page 149; 305pp; English.
                                                                                                                                                                                                                                                           Jarvis T,
                                                                                                                                                                                                              98US-0079678.
                                                                                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                         Roberts E,
                                                                                                                                                                                                                                                                               WPI; 1999-591315/50.
                                                                                                                       Homo sapiens
                                                                                                                                                                                                             27-MAR-1998;
                                                                                                                                            WO9950403-A2
                                                                                                                                                                                       24-MAR-1999;
                                                                                                                                                                 07-0CT-1999
                                                                                                                                                                                                                                                         Pavco PA,
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McSwiggen JA;

Coeshott C,

The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beca 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA1766 to AAA1762 represent ribozyme sequences for ARNT, corresponding target sequences; AAA1768 to AAA19185 and AAA19185 to AAA19185 and AAA19185 to AAA19185 and AAA19185 to AAA1189 to AAA2180 to AAA21 integrin subunit alpha-6, or integrin subunit beta-3 Sequence 17 BP; 5 A; 1 C; 4 G; 7 U; 0 other;

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ö
                                         Gaps
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0
          0.9%; Score 12.8; DB 1; Length 17;
50.0%; Pred. No. 2.7e+02;
ve 6; Mismatches 2; Indels
                                                                1178 TGTTTCTATAGGTGAG 1193
Query Match
Best Local Similarity 50.0%;
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AAX59454 standard; DNA; 17 BP. 2 udurudukakangka 17 RESULT 495 AAX59454 ID AAX5 XX

99US-0129390,

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Reconstituted human antibody; peptide antigen HM1.24; framework region; complementary determining region; CDR; anti-HM1.24 antibody; myeloma; humanised antibody; primer; 8s.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The specification describes a reconstituted human antibody recognizing the peptide antigen HM1.24. This human antibody contains natural human framework regions modified by amino acid substitutions to provide homogeneity with a previously designed framework region (which may arise from a human or non-human source); and complementary determining regions (CDR) derived from a non-human anti-HM1.24 antibody. The reconstituted antibody is useful in the treatment of diseases in which the surface antigen HM1.24 is implicated such as myeloma. The present sequence is used in the creation of the antibodies of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ribozyme; erythropoletin; granulocyte colony stimulating factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Reconstituted human antibody useful in the treatment of myeloma
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                                                                                                                   Primer used in construction of humanised anti-HM1.24 antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 12.8; DB 1;
Pred. No. 2.7e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 5 A; 7 C; 4 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 120; 256pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hammerhead ribozyme substrate #1082.
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nes 14, Conservative
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                                                                                                                                                                                                                                                                                                                                                              WO9918212-A1
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                                                                                                                                                                                                                                                                                                       Synthetic.
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Homo sapiens
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inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor
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0.9%; Score 12.8; DB 1;
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                                                   McSwiggen J;
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            (RIBO-) RIBOZYME PHARM INC
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Best Local Similarity
                                                   Zwick M,
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0; The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of exthropoletin, granulocyte colony stimulating factor protein and interferon alpha. Gaps . 0 Length 17; Indels AAF05405 standard; DNA; 17

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The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor. BAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interferon alpha.
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                                                                           McSwiggen J;
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                                                                                                                                                                                                                           Claim 18; Page 116; 164pp; English.
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                                  (RIBO-) RIBOZYME PHARM INC
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                                                                                                                              Ribozyme; erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss.
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AAC70618 standard; DNA; 17

Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; ss.

Single nucleotide polymorphism PCR primer #299,

(first entry)

09-FEB-2001

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAMT Displacement Protein (CDF). Inhibition of the repressors removes prevents inhibition of and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interferon alpha.
Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interferon alpha.
                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin -
                                                                                                                                                                                                                                                                                  Ribozyme; erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss.
                                                                                                   .
0
                                                                             Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; ive 0; Mismatches 2; Indels
                                                                                                 Indels
                                                                           ch 0.9%; Score 12.8; DB 1; I
1 Similarity 87.5%; Pred. No. 2.7e+02;
14; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Enzymatic and antisense nucleic acid inhibition useful for producing a granniant ---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 5 A; 6 C; 5 G; 1 T; 0 other;
                                                     BP; 6 A; 1 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                McSwiggen J;
                                                                                                                                                                                                                                                            Hammerhead ribozyme substrate #3379.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 54; Page 133; 164pp; English.
                                                                                                                     439 AGAAAGTTGCTGAAGT 454
                                                                                                                                                                                                 BP
                                                                                                                                         AGGAATTTGCTGAAGT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                               Pavco P,
                                                                                                                                                                                                                                                                                                                                                                               11-APR-2000; 2000WO-US09721.
                                                                                                                                                                                                                                                                                                                                                                                                     99US-0129390
                                                                                                                                                                                             AAF07122 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-647423/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Zwick M,
                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                      WO200061729-A2
                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                     12-APR-1999;
                                                       Sequence 17
                                                                                                                                                                                                                                        16-FEB-2001
                                                                                                                                                                                                                                                                                                                                                            19-0CT-2000.
                                                                                                                                                                                                                    AAF07122;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Blatt L,
                                                                            Query Match
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                                                                                      Best Loca
Matches
                                                                                                                                                                        RESULT 502
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Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis -

Claim 8; Fig 5; 214pp; English

Lander

Cargill M, Daley GQ, Ireland JS, Patil N, Sklar P;

WPI; 2000-611722/58.

Altshuler D, Lipshutz RJ,

(WHED ) WHITEHEAD INST BIOMEDICAL RES. (AFFY-) AFFYMETRIX INC.

99US-0127248.

31-MAR-1999;

30-MAR-2000; 2000WO-US08440.

WO200058519-A2

05-OCT-2000.

Homo sapiens

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The present invention is concerned with a number of human single mucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endoorine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                      .
0
                                                                                                                                                                                                                                                                      Score 12.8; DB 1; Length 17;
Pred. No. 2.7e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Single nucleotide polymorphism PCR primer #301.
                                                                                                                                                                                                                            Sequence 17 BP; 6 A; 7 C; 3 G; 1 T; 0 other;
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0
                                                                                                                                                                                                                                                                      0.9%;
                                                                                                                                                                                                                                                                                                                                                           259 CTCCTGGGCTGA 274
                                                                                                                                                                                                                                                                                                                                                                                         CICCIGGGIIGGGIGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC70621 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 14; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC70621;
                                                                                                                                                                                   diseases.
                                                                                                                                                                                                                                                                                                                                                                                                     16
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Gaps

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394 GCAGCAATGGCCCGGC 409

14; Conservative

Local Similarity

Best Loca Matches

1 GCAGAAATCGCCCGGC 16

g ð

RESULT 503 AAC70618/c

Bellon L;

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with a target sequence and contain at least one phosphoro(di)thicate link, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or receptor gene, are used to treat cancer (particularly of breast or receptor gene receptor. Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with a lerrations in phenotype, particularly for identification of therapeutic targets, and as research reagens (for RNA, in the same way that restriction endonucleases are used with DNA). The combination of modifications in (A) improves resistent concleases, binding affinity and/or activity. AAA23503 to AAA24747 represent oestrogen receptor hammerhead ribozyme sequences, and AAA26105 represent their corresponding target sequences. AAA2593 to AAA26105 represent ceptor his ribozyme sequences, and AAA26107 to AAA26108 represent their corresponding target sequences, AAA2503 to AAA26107 represent ceptor halphin ribozyme sequences, AAA26207 to AAAA26108 represent their corresponding target sequences, AAA26201 to AAAA2610 represent their characteristic corresponding target sequences. AAA26201 to AAAA2611 represent
                                                                                                                                                                                                                                            present invention describes nucleic acids (A) that interact stably a target sequence and contain at least one phosphoro(di)thioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1952.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonuclectide; gene expression modification; cancer; bosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          other ribozyme sequences and antimense oligonucleotides used in the exemplification of the present invention.
                                                                                                                                  New nucleic acids that interact, and optionally cleave, target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
Karpeisky A,
Haeberli P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Karpeisky A,
Haeberli P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.9%; Score 12.8; DB 1;
87.5%; Pred. No. 2.7e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 other;
Beigelman L, McSwiggen JA,
Wick M, Jarvis T, Woolf T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Beigelman L, McSwiggen JA,
Zwick M, Jarvis T, Woolf T,
                                                                                                                                                                                                  Claim 77; Page 79; 148pp; English.
                                                                                                                                                         sequences, used to treat cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1144 TTTTTTTTTTGGA 1159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
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Best Local Similarity 87,20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA25454 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JUL-2000 (first entry)
                          Zwick M,
                                                                                     WPI; 2000-013248/01
                       Reynolds M, Zwic)
Matulic-Adamic J;
Thompson JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thompson JD,
Reynolds M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  W09954459-A2
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23-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose used to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                                                                       Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothicate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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0
                                                                                                                                                                                                                                                                  Lander ES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ch 0.9%; Score 12.8; DB 1; Length 17; l Similarity 87.5%; Pred. No. 2.7e+02; 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                               Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 6 A; 7 C; 3 G; 1 T; 0 other;
                                                                                                                                                                                        (WHED ) WHITEHEAD INST BIOMEDICAL RES. (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                               Daley GO,
Sklar P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 8; Fig 5; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     274
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                                                                                                     30-MAR-2000; 2000WO-US08440
                                                                                                                                                  99US-0127248
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                259 CTCCTGGGCTGGCTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                           Cargill M,
Patil N, S
                                                                                                                                                                                                                                                                                                                             WPI; 2000-611722/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
              WO200058519-A2
                                                                                                                                                                                                                                                           Altshuler D,
Lipshutz RJ,
                                                                                                                                                31-MAR-1999;
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23-JUN-1998;
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                                                        05-0CT-2000
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Gaps

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Bellon L;

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with a target sequence and contain at least one phosphoro (di) thioate link, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or condentium), in vivo or by transforming cells ex vivo and implanting treated cells, or for other conditions associated with levels of cestrogen receptor. Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with argets, and as research reagents (for RNA, in the same way that restriction endonucleases are used with DNA). The combination of modifications in (A) improves resistance to nucleases, binding affinity and/or activity. AAA21503 to AAA4477 represent oestrogen receptor harmerhead ribozyme sequences, and AAA2593 to AAAA62107 to AAAA6218 represent their corresponding target sequences. AAA2593 to AAAA6219 to AAAA6218 represent their corresponding target sequences. AAA2593 to AAAA6219 to AAAA6218 represent their corresponding target sequences. AAA2503 to AAAA6219 to AAAA6218 represent their corresponding target sequences. AAA2503 to AAAA6219 to AAAA6211 represent exemptification of the modern exemples and antisense oligonucleotides used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOB; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolesia; VGTI; syndrome; APP; PSEN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                                                                                                   The present invention describes nucleic acids (A) that interact stably
                                                                                      New nucleic acids that interact, and optionally cleave, target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LDLR mutation correcting oligonucleotide SEQ ID NO: 3842.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             exemplification of the present invention.
                                                                                                                                                       Claim 77; Page 79; 148pp; English.
                                                                                                               sequences, used to treat cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1144 TTTTTTTTTTGGA 1159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABA80996 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 87.5
Matches 14; Conservative
                                            WPI; 2000-013248/01.
Matulic-Adamic J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antilipemic; ss
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The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CTFR, cyclin-dependent kinase inhibitor 2A (CDEMZA3, ARC, Factor V, Factor VIII, Factor IX, haemoglobin alpha locus apolipoprotein E (APOR). LDL receptor (LDLR), upp-glucuronosyltransferase (UGT1), amyloid precursor protein (APC), presenilin-1 (PSENI) and presenilin-2 (PSENI). These can be used in the gene therapy of diseases
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                                                                                                                                                                                                                                         Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterclaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention.
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                                                                                                                                       Rice MC
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30-OCT-2000; 2000US-244989P.
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Best Local Similarity 87.55,
Thes 14; Conservative
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                                                              (UYDE ) UNIV DELAWARE
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The present invention provides single-stranded oligomucleotides which can be used for the targeted alteration of genomic sequences, where the oligomucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, cetinoblastoma, BRCAL, BRCAS, CFTR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor V, Pactor VII, Factor IX, haemoglobin alpha locus (CDKN2A), APC, Factor V, Factor VII, Factor INDP, UnDP-glucuronosyltransferase (UGTI), amyloid precursor protein (APC), presentian-1 (PSEN1) and presentian-2 (PSEN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, happercholesterolaemia, thalassemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting
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oligonucleotides of the invention.
                                                                                                                                                                                                                                                                                    Claim 7; Page 252; 294pp; English.
                                                                                                            Rice MC;
01-JUN-2000; 2000US-208538P.
30-OCT-2000; 2000US-244989P.
                                                              (UYDE ) UNIV DELAWARE
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                                       0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; tive 0; Mismatches 2; Indels
Sequence 17 BP; 3 A; 7 C; 3 G; 4 T; 0 other;
                                  Query Match 0.9
Best Local Similarity 87.5
Matches 14; Conservative
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ABA81248 standard; DNA; 17 BP. (first entry) 24-JAN-2002 ABA81248; RESULT 509 

PSEN1 mutation correcting oligonucleotide SEQ ID NO: 4094.

Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MIH1; APOB; mismatch repair; MSH6; hyperlipidaemia; apolipoprotein B; LDIR; familial hypercholesterolosemia; MCTI; syndrome; APP; PSENI; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presenilin-1; Albheimmer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilipemic;

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WO200173002-A2.

04-OCT-2001

27-MAR-2001; 2001WO-US09761.

27-MAR-2000; 2000US-192176P.

27-MAR-2000; 2000US-192179P. 01-JUN-2000; 2000US-208538P. 30-OCT-2000; 2000US-244989P. 

(UYDE ) UNIV DELAWARE

Rice MC Gamper HB, Kmiec EB,

WPI; 2001-639230/73.

Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification -

Claim 7; Page 265; 294pp; English.

The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the cligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor VIII, Factor IX, haemoglobin alpha locus (CDKN2A), APC, Factor VIII, Factor IX, haemoglobin alpha locus (CDKN2A), APC, Factor VIII, Factor IX, haemoglobin alpha locus (UGTI), amyloid precursor protein (APC), presentlin-1 (PSENI) and presentlin-2 (PSEN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophila, hypercholesterolaemia, thalassaemia, sickle cell anaemia, various syndromes. The present sequence is one of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention.

Sequence 17 BP; 0 A; 2 C; 9 G; 6 T; 0 other;

Gaps .; 0 Length 17; Indels 0.9%; Score 12.8; DB 1; 17.5%; Pred. No. 2.7e+02; ve 0; Mismatches 2; 87.5%; Query Match Best Local Similarity 87.55 Matches 14; Conservative

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Gaps

ABA81249 standard; DNA; 17 RESULT 510 ABA81249,

BP.

24-JAN-2002 (first entry)

ABA81249;

PSEN1 mutation correcting oligonucleotide SEQ ID NO: 4095.

Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2, CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN24; melanoma; APC; HBA21; HBA2; adenomatous polyposis of the colon; Factor VI; Factor IX; Lhrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismacch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein B; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSEM1; antisense; UDP-glucuromosyltransferase; amyloid precursor protein; presentin; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilipemic; ss

Homo sapiens

WO200173002-A2

04-OCT-2001.

27-MAR-2001; 2001WO-US09761. 

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16-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic canonic at least one mismatch compared with the genomic conjourcleotide has at least one mismatch compared with the genomic canonic at least one at least one mismatch compared with the genomic canonic and present in particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, rection a CDKN2A), APC, Pactor VIII, Factor IX, haemoglobin alpha locus (CDKN2A), APC, Pactor VIII, Factor IX, haemoglobin alpha locus (CDKN2A), haemoglobin alpha locus 2 (HBA2), wiffit, WSH2, WSH6, apolipoprotein E (APOB), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGTI), amyloid precursor protein (APC), presentiln.1 (PSENI) and presentiln.2 (PSENI2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's diseases, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention.
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27-MAR-2000; 2000US-192176P.
27-MAR-2000; 2000US-192179P.
01-JUN-2000; 2000US-208538P.
                                                                        30-OCT-2000; 2000US-244989P
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Matches 14; Conserv
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ABK03155 standard; RNA; 17 (first entry) 12-MAR-2002 ABK03155; RESULT 511 THE REAL PROPERTY OF STATE OF

Human CD20 Inozyme #106.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; b=cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immundeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immuncytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; central nervous system injury; cheretypakonlar accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

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sapiens. Synthetic

WO200159103-A2.

ABK03627 standard; RNA; 17 BP.

RESULT 512 ABK03627/c ABK03627;

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regression of a neurite growth inhibitor gene (NOGO).

regulates expression of a neurite growth inhibitor gene (NOGO).

The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (aleaving RNA with a NNY molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NNY motif), a G-cleaver (cleaving RNA with a NNY motif) and amberzyme (cleaving RNA with a NNY motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably MG^2+. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting cuse of nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular NHL, wanther-cell lymphoma (NHL), cleaming a condition of immunodeficiency virus) associated NHL, manthe-cell lymphoma (NHL), immunodeficiency virus) associated NHL, manthe-cell lymphoma (MC), immunodeficiency virus) associated with a cell to reduce NOGO gene in the presence of a divalent cation that is preferably MG^2+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the cell and createment may further comprise the use of one or more therapies.

Contral nervous system (CNS) injury and created central nervode subset (CNS) injury and cell or or more therapies or cent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
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Pred. No. 2.7e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            and central nervous system injury
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Best Local Similarity 87.5%;
                                                                                     11-FEB-2000, 2000US-181797P.
28-FEB-2000, 2000US-185516P.
06-MAR-2000, 2000US-187128P.
                                             09-FEB-2001; 2001WO-US04273
                                                                                                                                                                                  RIBOZYME PHARM INC
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12-MAR-2002
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Human CD20 DNAzyme #81

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; incyme; G-Cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCI; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease, ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

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sapiens, Synthetic.

WO200159103-A2

16-AUG-2001

09-FEB-2001; 2001WO-US04273

28-FEB-2000; 2000US-185516P. 06-MAR-2000; 2000US-187128P. 11-FEB-2000; 2000US-181797P.

RIBOZYME PHARM INC. BLATT L. MCSWIGGEN J. (RIBO-) 1 (BLAT/) 1 (MCSW/) 1

CHOWRIRA B M.

Chowrira BM; McSwiggen J, Blatt L,

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury

Claim 30; Page 160; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down capression of a neutite growth inhibitor gene (NGGO).

The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (cleaving RNA with an NGN with an NGN with a NGN molecule ocid cleaving RNA with a YGY motif). The CD20-targetting nucleic acid is used (cleaving RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg^27+. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the cuse of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky converged or follicular NHL, lymphocytic leukaemia, Huphoma (MCL), immunocytoma (IMC), small B-cell lymphoca, immunocytoma (IMC), small B-cell lymphoca, immune confice acid is used to cleave RNA of the NGGO gene in the presence of a divalent cation that is preferably Mg^22+. Furthermore, the nucleic acid any be contacted with a cell to reduce NOGO activity of the cell and creatment may further comprise the use of one or more therapies.

Treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more therapies.

In particular, the NOGO-targetting nucleic acid may be used to treat central nervous system (CNS) injury and cerebrovascular accident CVS).

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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78752 to ABV78762 and AB989519 to ABB99520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was mapped to human chromosome 10912.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), Parkinson's disease, creutzfeldt-Jakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present sequence is a DNAzyme molecule of the invention.
                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL -
                                                                                                                                                                                                        0;
                                                                                                                                                                   Length 17;
                                                                                                                                                                 Score 12.8; DB 1; Length 1'
Pred. No. 2.7e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human HTPL scanning oligonucleotide SEQ ID 1816.
                                                                                                                              Sequence 17 BP; 4 A; 3 C; 2 G; 8 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Page 301; 718pp; English.
                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                             437 TCAGAAAGTTGCTGAA 452
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30-JAN-2001; 2001MO-US00665.
30-JAN-2001; 2001MO-US00667.
30-JAN-2001; 2001MO-US00669.
                                                                                                                                                                     0.98;
                                                                                                                                                                                      87.5%;
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09-OCT-2001; 2001US-0327898
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                                                                                                                                                                                                                                                                                                                                                                           ABV80570 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                              17 raagaaagrigereaa
                                                                                                                                                                                                        14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (AEOM-) AEOMICA INC
                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EP1229046-A2
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                                                                                                                                                                                                                                                                                                                                                                                                              ABV80570;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zhan J;
                                                                                                                                                                                                          Matches
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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, selectal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potential therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss. Gaps Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL -. 0 Length 17; Score 12.8; DB 1; Length 1 Pred. No. 2.7e+02; 0; Mismatches 2; Indels Human HTPL scanning oligonucleotide SEQ ID 1817 Sequence 17 BP; 5 A; 4 C; 4 G; 4 T; 0 other; Example 2; Page 302; 718pp; English. ..0 30-JAN-2001; 2001WO-US00664. 30-JAN-2001; 2001WO-US00665. 30-JAN-2001; 2001WO-US00667. 30-JAN-2001; 2001WO-US00669. 23-MAX-2001; 2001WO-US00669. 09-OCT-2001; 2001US-0864761. ВЪ 239 CATCTGCATCTGGGAC 254 0.9%; 28-JAN-2002; 2002EP-0001167. 17 CATGTTCATCTGGGAC 2 example from the invention. ABV80571 standard; DNA; 17 (first entry) 14; Conservative (AEOM-) AEOMICA INC WPI; 2002-676582/73 Local Similarity 30-JAN-2001; EP1229046-A2 Homo sapiens 03-JAN-2003 07-AUG-2002 ABV80571; Query Match Zhan J; Matches ABV80571,
TID AB 888666666888

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mapped to human chromosome 10p12.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or afternal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are calinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an example from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
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                                                                                                                                                                                                                                                                    0
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                                                                                                                                                                                                                                                                    Indels
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Pred. No. 2.7e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human HTPL scanning oligonucleotide SEQ ID 4084.
                                                                                                                                                                                                 Sequence 17 BP; 5 A; 4 C; 5 G; 3 T; 0 other;
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30-JAN-2001; 2001WO-US00669.
23-MAY-2001; 2001US-0864761.
09-OCT-2001; 2001US-0327898.
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Best Local Similarity 87.5°
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to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was mapped to human chromosome 10pl2.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an example from the invention.
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Score 12.8; DB 1; Length 17; Pred. No. 2.7e+02; 0; Mismatches 2; Indels Sequence 17 BP; 2 A; 4 C; 2 G; 9 T; 0 other; 1137 CTATGCCTTTTTTTT 1152 0.9%; Query Match
Best Local Similarity 87.5
Matches 14; Conservative

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Gaps

0;

crardcarrigrifici 17

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ABV82839 standard; DNA; 17 BP. ABV82839; 516 ABV82839 RESULT

(first entry) 03-JAN-2003

Human HTPL scanning oligonucleotide SEQ ID 4085.

Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis syrpessed Parched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.

Homo sapiens

EP1229046-A2

07-AUG-2002

28-JAN-2002; 2002EP-0001167

2001WO-US00664. 2001WO-US00665. 2001WO-US00663 30-JAN-2001; 30-JAN-2001;

2001WO-US00669. 2001US-0864761. 2001WO-US00667 2001WO-US00668 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 23-MAY-2001;

(AEOM-) AEOMICA INC

2001US-0327898

09-OCT-2001;

Zhan J;

WPI; 2002-676582/73.

Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL -

Example 2; Page 599; 718pp; English.

The present invention relates to human testis expressed Patched like protein (HTPL), see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL

·. shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was mapped to human chromosome 10pl2.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of such disorders associated with decreased expression or activity of human HTPL, Such disorders include disorders of testis, or adrenal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for example from the invention. The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino shares an overall structure organisation with the Patched protein. The Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss. Gaps Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -. 0 Length 17; Example 2; SEQ ID NO 220; 60pp + Sequence Listing; English. 2; Indels Human POSHL1 scanning oligonucleotide SEQ ID NO 220. 0.9%; Score 12.8; DB 1; 87.5%; Pred. No. 2.7e+02; Sequence 17 BP; 3 A; 3 C; 2 G; 9 T; 0 other; 0; Mismatches 1137 CTATGCCTTTTTTTT 1152 BP. 30-JAN-2001; 2001WO-US00664. 30-JAN-2001; 2001WO-US00665. 30-JAN-2001; 2001WO-US00666. 30-JAN-2001; 2001WO-US00667. 30-JAN-2001; 2001WO-US00668. 30-JAN-2001; 2001WO-US00669. 1 crardcarrrgrircr 16 2001WO-US00663 2001US-0864761 2001US-0328205 2002EP-0001165 ABV89507 standard; DNA; 17 (first entry) 14; Conservative WPI; 2002-684061/74. (AEOM-) AEOMICA INC Best Local Similarity 28-JAN-2002; 30-JAN-2001; Homo sapiens. EP1239051-A2. 23-MAY-2001; 10-OCT-2001; 23-DEC-2002 11-SEP-2002 Shannon M; ABV89507; Query Match Matches RESULT 517 ABV89507 888888888888888888888888888888888 6

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acids (S1, ABBB3999), a sequence having 65% sequence identity to (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and mucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarraps which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The
                                                                                                                                                                                                                     present sequence is that of a scanning oligonucleotide useful in examples of the invention.
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Rho GTPase; signal transduction; gene expression; cancer; vaccine;
gene therapy; transgenic; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human POSHL1 scanning oligonucleotide SEQ ID NO 221.
                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 2 A; 7 C; 4 G; 4 T; 0 other;
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                                                                                                                                                                                                                                                                                                the European Patent Office.
                                                                                                                                                                                                                                                                                                                                                                                                                                  160 CGCTGATCCTCAAGGT 175
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2001WO-US00665.
2001WO-US00666.
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Human POSHL 1 is a proto-congene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acide (II) encoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHLI including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic onn-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention.
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                              The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABB83999), a sequence having 65% sequence identity to (SI), (SI) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids.
                                                                                                                                                                                                                                                                                                                                             Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to Derwent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;
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Example 2; SEQ ID NO 221; 60pp + Sequence Listing; English.
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30-JAN-2001;
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SL4, ABBR399), a sequence having 65% sequence identity to (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small Grpases as well as downstream components of the signal transduction pathway. (I) is useful seconding (I) are useful for flagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and treating caused by altered expression of human POSHL1 including diagnosing and treating caused by electric and for surveying gene expression and creating care useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples
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Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
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                                                                         Example 2; SEQ ID NO 1247; 60pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 4 A; 9 C; 2 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the European Patent Office.
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2001WO-US00667. 2001WO-US00668. 2001WO-US00669.

30-JAN-2001;

30-JAN-2001; 23-MAY-2001; 10-OCT-2001;

30-JAN-2001; 30-JAN-2001; 2001WO-US00670. 2001US-0864761. 2001US-0328205.

(AEOM-) AEOMICA INC

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino acids (S1, ABBR399), a sequence having 65% sequence identify to (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous anino acids. Human POSHL 1 is a proto-oncogene product that functions as an adaptor protein that interacts with Rho family small Grpases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) for identifying a specific binding partner. (I) and nucleic acids (II) consoding (I) are useful for diagnosing, monitoring disease and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-numan animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                    Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
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                                                                                                                                                Example 2; SEQ ID NO 1248; 60pp + Sequence Listing; English.
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87.5%; Pred. No. 2.7e+02;
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30-JAN-2001; 2001WO-US00665.
30-JAN-2001; 2001WO-US00666.
30-JAN-2001; 2001WO-US00666.
30-JAN-2001; 2001WO-US00669.
30-JAN-2001; 2001WO-US00669.
23-MAY-2001; 2001WO-US00670.
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Best Local Similarity 87.5.
Best Local 14; Conservative
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                                     WPI; 2002-684061/74.
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ABV91053
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                                                                                                                                                             The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABBB3999), a sequence having 65% sequence of 6710, (SI), having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying aspecific binding partner. (I) and mucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating cancer, they useful in the development of waccines and (II) is useful for measuring and for surveying gene expression and creating are useful for measuring and for surveying gene expression and creating transgent onon-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples
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                                                                                           Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
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                                                                                                                                          Example 2; SEQ ID NO 1259; 60pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             268 TGGCTGATCAAAGAGG 283
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2001WO-US00665.
2001WO-US00666.
10-OCT-2001; 2001US-0328205
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                      (AEOM-) AEOMICA INC
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Best Local Similarity
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30-JAN-2001; 2
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30-JAN-2001; 2
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino acids (S1, ABBB3399), a sequence having 65% sequence identity to (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPasses as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) and coding (I) are useful for diagnosing, monitoring disease and traating cauced by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention.
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30-JAN-2001; 2001WO-US00668.
30-JAN-2001; 2001WO-US00669.
30-JAN-2001; 2001WO-US00670.
23-MAY-2001; 2001US-0864761.
10-OCT-2001; 2001US-0328205.
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nes 14; Conservative
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                                                                                                                                                                                                                                                                                                             Shannon M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABV91053;
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2001WO-US00664. 2001WO-US00665. 2001WO-US00666. 2001WO-US00667. 2001WO-US00668.

2001WO-US00670. 2001US-0864761. 2001US-0328205.

2001WO-US00669.

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (G1, ABBB3999), a sequence having 65% sequence identity to (S1), (S1), having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) consetul grancer, they useful in the development of vaccines and treating caused by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; SEQ ID NO 1767; 60pp + Sequence Listing; English.
                        28-JAN-2002; 2002EP-0001165
                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-684061/74.
                                                                                                                                                                                                                                                                                                                                                         (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention.
                                                                                             30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                   10-OCT-2001;
                                                                                                                                                                                                                                                   30-JAN-2001;
                                                                                                                                                                                                                                                                                23-MAY-2001;
                                                                        30-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                         Shannon M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of the
g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (S1, ABBB3939), a sequence having 55% sequence identity to (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) for identifying a specific binding partner. (I) and nucleic acids (II) caused by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to Derwent by the European Patent Office.
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                                                                                                                                                                                                                                                                                                                                                                                              Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; SEQ ID NO 1766; 60pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 5 A; 3 C; 5 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABV91054 standard; DNA; 17 BP.
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                                                                                             30-JAN-2001; 2001WO-US00668.
30-JAN-2001; 2001WO-US00669.
30-JAN-2001; 2001WO-US00670.
23-MAY-2001; 2001US-0864761.
10-OCT-2001; 2001US-0328205.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 87.5%;
les 14; Conservative
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                        2001WO-US00665.
2001WO-US00666.
2001WO-US00667.
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                                                                                                                                                                                                                                                      (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention.
                        30-JAN-2001;
                                                                     30-JAN-2001;
                                                                                                                                                                                                                                                                                                           Shannon M;
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Matches
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NAMES OF COLOR COL

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Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to Derwent by the European Patent Office.
                                                                                                                                              Gaps
                                                                                                                                                ..
                                                                                                          Length 17;
                                                                                                            Score 12.8; DB 1; Length 1'
Pred. No. 2.7e+02;
0; Mismatches 2; Indels
                                                                            Sequence 17 BP; 4 A; 3 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                          Human KTOMla portion (ABQ63232) probe # 347.
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0
                                                                                                                                                                                    906 GGCCCTGGTCCTAAAG 921
                                                                                                              0.9%;
                                                                                                                                                                                                                    16 GACCCTTGTCCTAAAG 1
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                                                                                                              Query Match 0.9
Best Local Similarity 87.5
Matches 14; Conservative
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                                                                                                                                                                                                                                                                             RESULT 525
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Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GFPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss.

EP1239051-A2

Ното

BXBXBXBXBXBXBXB

11-SEP-2002

Human POSHL1 scanning oligonucleotide SEQ ID NO 1767.

(first entry)

23-DEC-2002

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2001WO-US00661. 2001WO-US00662. 2001WO-US00663.

2000US-236359P. 2000GB-0024263.

2001WO-US00666. 2001WO-US00667. 2001WO-US00668.

2001WO-US00669. 2001US-0864761. 2001WO-US00670

2001WO-US00664. 2001WO-US00665.

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21-SEP-2001; 2001WO-US29656.
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                     WO200224750-A2
    Homo sapiens.
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30-JAN-2001;
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                                                                               21-SEP-2000;
                                                                                                  04-OCT-2000;
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                                                                                                                                                                                                                                                                                                                                                                  KTOM1 (kidney tumour overexpressed membrane) protein. The protein of the invention has cyrostatic activity. The nucleotide may have a use in gene therapy. The KTOM1 nucleic actions may be used to diagnose, treat or monitor a disease caused by altered expression of human KTOM1.

Compositions comprising the nucleic action, proteins or antibodies may be used to treat subjects having defects in KTOM1 which can manifest may be early of the kidney, as well as a disorder of liver, bone marrow, brain, heart, lung, kidney, as well as a probe used in the invention to scance the requence represents a probe used in the invention to
                                                                                                                                                                                                                                                                                   New human kidney tumor overexpressed membrane (KTOM1) protein and nucleic acids encoding the protein, useful for treating subjects having defects in KTOM1 which can manifest as cancer of the kidney, or as a disorder of e.g., liver or bone
                                                                                                                                                                                                                                                                                                                                                          invention relates to a novel isolated nucleic acid encoding human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ..
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 2 A; 8 C; 5 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                      Example 2; Page 203; 418pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1249 GCCATGTGAGGCCAGG 1264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABQ63636 standard; DNA; 17 BP
                                                                                              2001WO-US00661,
2001WO-US00662,
                                                                                                                  2001WO-US00663,
2001WO-US00664,
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                                                                                                                                                         30-JAN-2001; 2001WO-US00667.
                                                                                                                                                                  2001WO-US00668.
2001WO-US00669.
                                                                                                                                                                                      2001WO-US00670.
2001US-0864761.
                                               21-SEP-2001; 2001WO-US29656
                                                                                     000GB-0024263
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                                                                                                                                                                                                                            (AEOM-) AEOMICA INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
          WO200224750-A2
                                                                                                                 30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                              30-JAN-2001;
30-JAN-2001;
                                                                                                                                               30-JAN-2001;
                                                                                      04-OCT-2000;
                                                                                                                                                                             30-JAN-2001;
                                                                            27-SEP-2000;
                            28-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14;
                                                                                                                                                                                                                                                Zhang J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 526
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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The invention relates to a novel isolated nucleic acid encoding human kTCM1 (kidney tumour overexpressed membrane) protein. The protein of the invention has cytostatic activity. The nucleotide may have a use in gene therapy. The KTOM1 nucleotide may he used to diagnose, treat or monitor a disease caused by altered expression of human KTOM1.

Compositions comprising the nucleic acids, proteins or antibodies may be used to treat subjects having defects in KTOM1 which can manifest as cancer of the kidney, as well as a disorder of liver, bone marrow, brain, heart, lung, kidney, colon, skeletal muscle, testis, uterus and placenta function. The sequence represents a probe used in the invention to scan the nt 1-1001 portion of human KTOM1a (ABQ63232).
                                                                                                                                                                     New human kidney tumor overexpressed membrane (KTOM1) protein and nucleic acids encoding the protein, useful for treating subjects having defects in KTOM1 which can manifest as cancer of the kidney, or as a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, chloride channel calcium activated 1; CLCA1; ss; antiasthmatic; antiinflammatory; chronic obstructive pulmonary disease; COPD, asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 2 A; 9 C; 5 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                               Example 2; Page 203; 418pp; English.
                                                                                                                                                                                                                                                                    disorder of e.g., liver or bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1248 GGCCATGTGAGGCCAG 1263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABK55869 standard; RNA; 17 BP.
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                                                                                                                    WPI; 2002-479509/51.
(AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK55869;
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                                                             Zhang J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Loca
Matches
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Human, KTOMla, KTOMl; kidney tumour overexpressed membrane, cytostatic, gene therapy, cancer, kidney, liver, bone marrow, brain, heart, lung, kidney, colon, skeletal muscle, testis, uterus, placenta, probe, ss.

Human KTOMla portion (ABQ63232) probe # 349.

(first entry)

20-AUG-2002

ABQ63636;

ö

WO200211674-A2.

Homo sapiens.

therapy; bronchodilator; corticosteroid; vaccination; mucokinetic;

oxygen

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The invention relates to enzymatic nucleic acid molecules that down regulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic obstructive pulmonary disease (COPD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions that are related to will respond to the levels of CLCA1 in a cell or tissue. The sequences are useful for reducing CLCA1 activity in a cell, hence, are useful for treatment of a patient having a condition associated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapies under conditions antiable for the antibacterials, vaccinations, acetyloysteine and mucokinetic agents. The nucleic acids of the invention are also used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an
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                                                                                                                                                                                                                                                                                                                                                                                                               Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                    Szymkowski DE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.9%; Score 12.8; DB 1; Length 17; 68.8%; Pred. No. 2.7e+02; rative 3; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                    Ayers D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human CLCAl gene enzymatic nucleic acid #416.
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                                                                                                                                                                                                                                                                                                                    McSwiggen J, McKenzie T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; Page 57; 152pp; English.
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                                                                                                                                                                    09-AUG-2001; 2001WO-US24970.
                                                                                                                                                                                                         09-AUG-2000; 2000US-224383P.
                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC.
(SYNT ) SYNTEX USA LLC.
(THOM/) THOMPSON J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acetylcysteine
                                                                                            WO200211674-A2
                 acetylcysteine
                                                       Homo sapiens.
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                                                                                                                                                                                                                                                                                                                    Thompson J,
                                                                                                                              14-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                          Grupe A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 528
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The invention relates to enzymatic nucleic acid molecules that down regulate expression of chloride channel calcium activated I (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic obstructive pulmonary disease (COPD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowell syndrome and any other diseases or conditions that are related to or will respond to the levels of CLCA1 in a cell, there, are useful for treatment of a patient having a condition associated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, acetyloysteine and mucokinetic agents. The nucleic acids of the invention are also used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Szymkowski DE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Thompson J, McSwiggen J, McKenzie T, Ayers D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.9%; Score 12.8; DB 1; 17.5%; Pred. No. 2.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human CLCAl gene enzymatic nucleic acid #991.
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); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; Page 60; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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                                                                                                                                                                                                      09-AUG-2001; 2001WO-US24970.
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                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM
                                                                                                                                                                                                                                                                                                                                                                         (SYNT ) SYNTEX USA LLC. (THOM/) THOMPSON J.
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                                                                                                                                   14-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Grupe A;
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SXEXEXEXEXER EXECUTION OF SOCIOUS SECTION OF SOCIOU
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Page 259

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The invention relates to enzymatic nucleic acid molecules that down regulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic obstructive pulmonary disease (COPD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions that are related to will respond to the levels of CLCA1 in a cell or tissue. The sequences are useful for reducing CLCA1 activity in a cell, hence, are useful for treatment of a patient having a condition associated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapies under conditions suitable for the artibacterials, vaccinations, acetyloysteine and mucokinetic agents. The nucleic acids of the invention are also used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an
                                                                                                                                                                                                                                                             Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human CLCA1 gene enzymatic nucleic acid #992.
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                                                                                                                                                                               McKenzie T,
                                                                                                                                                                                                                                                                                                                              Claim 4; Page 76; 152pp; English.
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                                              09-AUG-2001; 2001WO-US24970
                                                                             09-AUG-2000; 2000US-224383P
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                                                                                                              (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                               Thompson J, McSwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                               USA LLC
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                                                                                                                                              THOM/) THOMPSON J.
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tes 14; Conserv
                                                                                                                               SYNTEX
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                 14-FEB-2002.
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                                                                                                                                                                                                Grupe A;
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Matches
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                                     Gaps
                                     0;
0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; Live 0; Mismatches 2; Indels
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Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic; antiinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinetic;

Homo sapiens.

WO200211674-A2

14-FEB-2002

09-AUG-2001; 2001WO-US24970.

09-AUG-2000; 2000US-224383P.

(RIBO-) RIBOZYME PHARM (SYNT ) SYNTEX USA LLC.

Szymkowski DE; Ayers D, McKenzie T, McSwiggen J, Thompson J, Grupe A;

(THOM/) THOMPSON J.

Szymkowski

Ayers D,

Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma WPI; 2002-217145/27.

Claim 4; Page 76; 152pp; English.

The invention relates to enzymatic nucleic acid molecules that down regulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic obstructive pulmonary disease (COPD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions that are related to or will respond to the levels of CLCA1 in a cell or issue. The sequences are useful for reducing CLCA1 activity in a cell, hence, are useful for treatment of a patient having a condition associated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, exygen therapy, bronchdilators, corputiositions antibacterials, vaccinations, acetylcysteine and mucokinetic agents. The nucleic acids of the invention are also used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an

Sequence 17 BP; 7 A; 5 C; 2 G; 3 U; 0 other;

Gaps ö Length 17; 0.9%; Score 12.8; DB 1; Length 1 87.5%; Pred. No. 2.7e+02; vative 0; Mismatches 2; Indels 14; Conservative Query Match Best Local Similarity Matches

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1149 TTCTTTTGGAAGTAA 1164 rrcerrrecaacica 16

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ABK57130 standard; RNA; 17 BP.

ABK57130;

02-JUL-2002 (first entry)

Human CLCAl gene enzymatic nucleic acid #1501.

oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinetic; Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic; antiinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; acetylcysteine 

Homo sapiens,

WO200211674-A2.

14-FEB-2002.

09-AUG-2001; 2001WO-US24970

09-AUG-2000; 2000US-224383P.

Chronic obstructive

Enzymatic polynucleotide that down regulates expression of channel calcium activated gene, useful for treating Chronic pulmonary disease (COPD), chronic bronchitis and asthma -

Claim 4; Page 112; 152pp; English.

chloride

Szymkowski DE;

McKenzie T, Ayers D,

Thompson J, McSwiggen J,

Grupe A;

(THOM/) THOMPSON J.

WPI; 2002-217145/27.

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Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic; antiinflammatory; chronic obstructive pulmonary disease, COPP, asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinetic;
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                                                                                                                                                                                                                                                                                                                                                                                                       ABK57404 standard; RNA; 17 BP
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Best Local Similarity 68.8%;
Matches 11; Conservative
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      (RIBO-) RIBOZYME PHARM INC.
(SYNT ) SYNTEX USA LLC.
(THOM/) THOMPSON J.
                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                 WPI; 2002-217145/27.
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                                        Thompson J,
                                                                                                                                                                                                                                                                                                                                                                                                                       ABK57404;
                                                 Grupe A;
                                                                                                                                                                                                                                                                                                                                                                                      532
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ABK57404/
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The invention relates to enzymatic nucleic acid molecules that down cregulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic constructive pulmonary disease (CPD), chronic bronchitis, asthma, cystic chart are related to or will respond to the levels of CLCA1 in a cell or tissue. The sequences are useful for reducing CLCA1 activity in a cell, hence, are useful for treatment of a patient having a condition consisted with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the associated with the level of CLCA1, where the invention further comprises treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, acetylcysteine and mucokinetic agents. The nucleic acids of the invention are also used as diagnostic tools to camine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an exympatic nucleic acid molecule of the invention.
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                                                                                                                                                                                                                                                                                                                         Enzymatic polynucleotide that down regulates expression of chloride
                                                                                                                                                           Szymkowski DE;
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8.8%; Pred. No. 2.7e+02;
ve 3; Mismatches 2; Indels
                                                                                                                                                                 McSwiggen J, McKenzie T, Ayers D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 4 A; 3 C; 6 G; 4 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 96; 152pp; English.
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                                                                                                                                                                Gaps
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                                                                          Length 17;
                                                                                                                                                            Indels
                                                                          0.9%; Score 12.8; DB 1; 37.5%; Pred. No. 2.7e+02; ve 0; Mismatches 2;
Sequence 17 BP; 8 A; 5 C; 2 G; 2 U; 0 other;
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2001WO-US00664.
2001WO-US00665.
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30-JAN-2001; 2001WO-US00662.
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2000US-234687P.
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                                                                                                                           87.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABN01790 standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                         16 rrgrrcgrrrrggad
                                                                                                                           Best Local Similarity 87.5
Matches 14; Conservative
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30-JAN-2001;
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27-SEP-2000;
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification and substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise concentration and/or amount specifically of hGDMLP proteins as specific biomolecule capture probes for surface-enhanced laser desorption concentration, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement the arrapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders on hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1 -
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                                                                                                                                                                                                                            Chen W, Shannon ME;
                                                                                                                                                                                                                                                                                                                  New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1
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Shannon ME;

Chen W,

Rank DR,

Hanzel DK,

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N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
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                                         ftp.wipo.int/pub/published_pct_sequence.
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WO200192524-A2 Homo sapiens.

RESULT 534 ABN01792

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0.9%; Score 12.8; DB 1; Length 17;

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                                                                                                                                    Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                     Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1889.
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 mucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and cor expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concration and/or amount specifically of hGDMLP proteins, as specific blomclecule capture probes for surface-enhanced laser desorption constantion, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement
                                                                                                                     Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; 88.
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                                                                                      Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1890.
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therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                 N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                                                  0.9%; Score 12.8; DB 1;
ilarity 87.5%; Pred. No. 2.7e+02;
Conservative 0; Mismatches 2;
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                                                                                                                                            Sequence 17 BP; 2 A; 5 C; 7 G; 3 T; 0 other;
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30-JAN-2001; 2001WO-US00670.
05-FEB-2001; 2001US-266860P.
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27-SEP-2000;
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                                                                           invention.
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protein 1 (hGDWLP-1). The protein and polymucleotide sequences of hGDWLP-1 can be used in gene therapy and vaccine production. The hGDWLP-1 mucleic acids can be used as probes to derect, characterise and quantify hGDWLP-1 mucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDWLP-1 protein variants having desired phenotypic improvements, and correstration and/or amount specifically proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDWLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDWLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption in action, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders hGDMLP-1, in the part and skeletal muscle disorders hGDMLP-1, in the part and skeletal muscle disorders.
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87.5%; Pred. No. 2.7e+02;
rative 0; Mismatches 2;
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30-JAN-2001; 2001WO-US00667
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Best Local Similarity
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(AEOM-) AEOMICA INC

Shannon ME;

Chen W,

Rank DR,

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of brotein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used as probes to detect, characterise and quantify hGDMLP-1 mucleic acids in samples, as amplification and quantify hGDMLP-1 mucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise concentration and/or amount specifically of hGDMLP proteins, as specific concentration, as therapeutic supplement in patients having specific concentration, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in not particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence in the exemplification of the present
                                                                                                          New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1
                                                                                                                                                                                                                                    Disclosure, SEQ ID 6159; 214pp; English.
                   Hanzel DK,
                                                              WPI; 2002-179446/23.
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                      Gu Y,
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N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO Gaps . 0 0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; tive 0; Mismatches 2; Indels Sequence 17 BP; 2 A; 7 C; 5 G; 3 T; 0 other; ftp.wipo.int/pub/published\_pct\_sequence. Query Match Best Local Similarity 87.5; Matches 14; Conservative

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257 ACCICCIGGGCIGGCI 272 1 ACCTCACGGCTGGCT 16 à

RESULT 539

ABN06274 standard; DNA; 17 BP. ABN06274; 

(first entry) 29-MAY-2002

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6266.

Homo sapiens.

WO200192524-A2.

06-DEC-2001

; 2000US-207456P. ; 2000US-2346B7P. ; 2000US-236359P. ; 2000GB-0024263. 25-MAY-2001; 2001WO-US16981. 21-SEP-2000; 27-SEP-2000; 04-OCT-2000; 26-MAY-2000;

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 proteins variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise (C for expressing the proteins. The hGDMLP-1 proteins as specific concentration and/or amount specifically of hGDMLP proteins, as specific concentration, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement configuration, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in continue the present sequence represents an oligomer used in the present in the exemplification of the present in the exemplification of the present N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequence. Shannon ME; New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human Chen W, Hanzel DK, Rank DR, Disclosure; SEQ ID 6266; 214pp; English. myosin-like protein hGDMLP-1 2001WO-US00668. 2001WO-US00669. 2001WO-US00665. 2001WO-US00662. 2001WO-US00663. 2001WO-US00664 2001WO-US00666 2001WO-US00670 05-FEB-2001; 2001US-266860P Penn SG, WPI; 2002-179446/23. (AEOM-) AEOMICA INC. 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; Ji Y, Gu Y, 

Gaps 0 0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; cive 0; Mismatches 2; Indels Conservative Local Similarity 14; Query Match Matches

Sequence 17 BP; 2 A; 8 C; 5 G; 2 T; 0 other;

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371 GGGCCCAGCTTCCTCC 386

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ABN06275 standard; DNA; 17 BP dedeceadecrecerce 17 (first entry) 29-MAY-2002 ABN06275; RESULT 540 ABN06275

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6267.

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

us09904568-3.rng

RESULT 541

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymuclectie sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 mucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hODMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specifically of hGDMLP proteins, as specifically of hGDMLP proteins, as specifically editions, as therapeutic supplement in patience having specific there is a standard of a surface-enhanced laser desorption is a characteric supplement in patience having specific there is a specific and a series of the production, and in vaccines or for replacement and the production, and in vaccines or for replacement and the production and 
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                                Homo sapiens.
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Shannon ME;

Chen W,

Rank DR,

Hanzel DK,

Penn SG,

Ji Y,

2001WO-US00670 2001US-266860P

2001WO-US00663. 2001WO-US00664. 001WO-US00666

2001WO-US00665 2001WO-US00667 2001WO-US00668 2001WO-US00669

2000US-236359P 2000GB-0024263 2001WO-US00661 2001WO-US00662

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise CC chGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption concentration, an therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for therapy. The polynucleotide sequences encoding hGDMLP-1 in patients associated with the expression of hGDMLP-1, in particular heart and skelteral muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the present incomenting of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                               Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                              Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6507.
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              ABN06515 standard; DNA; 17 BP.
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27-SEP-2000;
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                                                 ABN06515;
ABN06515
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N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO

invention.

Gaps

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0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; tive 0; Mismatches 2; Indels

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14; Conservative

Local Similarity

Query Match

Best Loca Matches

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Gaps

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Query Match

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hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for particular heart and skeletal muscle disorders. hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                            N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence.
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Pred. No. 2.7e+02;
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30-JAN-2001; 2001WO-US00664.
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2001US-266860P.
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Best Local Similarity 87.5%;
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             The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vacaine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise
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                                                                                        0.9%; Score 12.8; DB 1;
87.5%; Pred. No. 2.7e+02;
tive 0; Mismatches 2;
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                                                     Sequence 17 BP; 5 A; 6 C; 4 G; 2 T; 0 other;
             at ftp.wipo.int/pub/published_pct_sequence.
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Shannon ME;

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                                                                                                                                                                                                                                                                                                              N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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                          Disclosure; SEQ ID 8381; 214pp; English.
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 proteins thaving desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise concentration and/or amount specifically of hGDMLP proteins, as specific concentration, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement the concentration, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement the rarpy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in no particular heart and skeletal muscle disorders and of hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of brOMMLP-1 can be used in gene therapy and vaccine production. The hCDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically proteins, as specific concentration, as therapeutic supplement in patients having specific concentration, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders subsome 12. The present sequence in the exemplification of the present
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2001WO-US00670.
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Gaps ; 0 0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; ative 0; Mismatches 2; Indels Sequence 17 BP; 5 A; 3 C; 7 G; 2 T; 0 other; Best Local Similarity Matches 14; Conserv Query Match

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29-MAY-2002 (first entry)

BP. ABN09588 standard; DNA; 17 ABN09588; 17 RESULT 546 ABN09588/c g

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:9580. 30-JAN-2001; 2001WO-US00663. 30-JAN-2001; 2001WO-US00664. 30-JAN-2001; 2001WO-US00665. 30-JAN-2001; 2001WO-US00666. 30-JAN-2001; 2001WO-US00667. 30-JAN-2001; 2001WO-US00668. 25-MAY-2001; 2001WO-US16981 2000US-236359P 2000GB-0024263 2001WO-US00661 30-JAN-2001; 2001WO-US00662 2001US-266860P WO200192524-A2 30-JAN-2001; Homo sapiens 27-SEP-2000; 34-OCT-2000; 36-DEC-2001 

Shannon ME;

Chen W,

3 Chen DR, Rank DK, Hanzel Gu Y, Ji Y, Penn SG, WPI; 2002-179446/23.

(AEOM-) AEOMICA INC.

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1

Disclosure; SEQ ID 9580; 214pp; English.

The present invention describes a human genome-derived myosin-like hopotein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification enubstrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present

N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequence.

Sequence 17 BP, 5 A, 3 C, 7 G, 2 T, 0 other;

Gaps .. 0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; ative 0; Mismatches 2; Indels 14; Conservative Query Match Best Local Similarity Matches 14; Conserv

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The present invention describes a human genome-derived myosin-like protein 1 (hGDWLP-1). The protein and polynuclectide sequences of hGDWLP-1 can be used in gene therapy and vaccine production. The hGDWLP-1 uncleic acids can be used as probes to detect, characterise and quantify hGDWLP-1 mucleic acids in samples, as amplification and quantify hGDWLP-1 mucleic acids in samples, as amplification of bGDWLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDWLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDWLP-1 proteins, as standards in assays used to determine the hGDWLP-1 proteins, as standards in assays used to determine the hGDWLP-1 proteins, as therapeutic supplement in patients having specific biomolecule capture probes for surface-enhanced laser desorption instain, as therapeutic supplement in patients having specific deficiency in hGDWLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDWLP-1 may be used for theraposing a disorder associated with the expression of hGDWLP-1, in particular heart and skeletal muscle disorders. HGDWLP-1 is localised to
                                                                                                                                                                                                                                  Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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Shannon ME;

Chen W,

Rank DR,

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chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                     N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence.
                                                                                                                                                                                                                                                                                                                                                                                     Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; 88.
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37.5%; Pred. No. 2.7e+02;
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                                                                                                                                                                                                                                                                                   N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence.
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as amplification
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87.5%; Pred. No. 2.7e+02;
tive 0; Mismatches 2; Indels
and quantify hGDMLP-1 nucleic acids in samples,
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 mucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption of hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
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                                                                New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface enhanced laser desorption ionization, comprises human
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WPI; 2002-179446/23.
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 mucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification cubstrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific blomolecule capture probes for surface-enhanced laser desorption concentration, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence in the exemplification of the present sequence in the exemplification of the present
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                                                             30-JAN-2001; 2001WO-US00667.
30-JAN-2001; 2001WO-US00668.
30-JAN-2001; 2001WO-US00669.
30-JAN-2001; 2001WO-US00670.
30-JAN-2001; 2001WO-US00665.
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Shannon ME;

Chen W,

Rank DR,

Hanzel DK,

Penn SG,

Gu Y, Ji Y,

WPI; 2002-179446/23.

2001WO-US00666.

2001WO-US00663. 2001WO-US00665

30-JAN-2001; 30-JAN-2001;

30-JAN-2001; 27-SEP-2000;

2000US-236359P 2000GB-0024263 2001WO-US00661 2001WO-US00662 2001WO-US00664

25-MAY-2001; 2001WO-US16981

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30-JAN-2001; 2001WO-US00667. 30-JAN-2001; 2001WO-US00668. 30-JAN-2001; 2001WO-US00669. 30-JAN-2001; 2001WO-US00669.

30-JAN-2001; 30-JAN-2001; 30-JAN-2001;

05-FEB-2001; 2001US-266860P

(AEOM-) AEOMICA INC.

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific blomolecule capture probes for surface-enhanced laser description ionization, comprises human myosin-like protein hGDMLP-1

Disclosure; SEQ ID 10727; 214pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in patiental heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
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Gaps

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0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; ve 0; Mismatches 2; Indels

Query Match 0.9%; Best Local Similarity 87.5%; Matches 14; Conservative

257 ACCTCCTGGGCTGGCT 272

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1 Accrecirecirecient 16

ABN10735 standard; DNA; 17 BP

RESULT 55 ABN10735/

RESULT 552 ABN10736/c ID ABN10736 standard; DNA; 17 BP.

Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:10727.

(first entry)

29-MAY-2002

ABN10735;

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WPI; 2002-179446/23.
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Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease;
                                                                            Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:10728.
                                                                                                                                     muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
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2001WO-US00665.
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                                      (first entry)
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30-JAN-2001;
30-JAN-2001;
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ABN10736;
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Hanzel DK, Rank DR, Gu Y, Ji Y, Penn SG, (AEOM-) AEOMICA INC.

Shannon ME;

Chen W,

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific blomolecule capture probes for surface-enhanced laeer desorption ionization, comprises human myosin-like protein hGDMLP-1

Disclosure; SEQ ID 10728; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of bGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 mucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific concentration, as therapeutic supplement in patients having specific concentration, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. HGDMLP-1 is localised to chromosome 22. The present sequence in the exemplification of the present contents of the hGDMLP-1 sequence in the exemplification of the present N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO

Sequence 17 BP; 1 A; 5 C; 8 G; 3 T; 0 other;

ftp.wipo.int/pub/published\_pct\_sequence.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption

Chen W, Shannon ME;

Rank DR,

Hanzel DK,

Gu Y, Ji Y, Penn SG, WPI; 2002-179446/23.

(AEOM-) AEOMICA INC.

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1 -

Disclosure; SEQ ID 10729; 214pp; English.

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                                                                                                                                                                Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                      Gaps
                                                                                                                                                 Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:10729.
                      ö
      Length 17;
                      Indels
      Score 12.8; DB 1;
Pred. No. 2.7e+02;
                       0; Mismatches
                                                                                                 ABN10737 standard; DNA; 17 BP.
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2001WO-US00668.
2001WO-US00669.
2001WO-US00670.
      0.9%;
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30-JAN-2001; 2001WO-US00663.
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                                                        16 ggagcccagcarccc 1
                                                                                                                                  (first entry)
Query Match
Best Local Similarity 87.5%
Thes 14; Conservative
                                       636 GGAGCTCTGCATCCCC
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30-JAN-2001; 2
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04-OCT-2000;
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ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucloctide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                 The sequence data for this patent did not form part of the printed lification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart, muscle, myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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0.9%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                           specification, but was obtained in electronate ftp.wipo.int/pub/published_pct_sequence.
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                                                                                                     invention.
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynuclectide sequences of hGDMLP-1 and be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description
                                                                                                                                                                                                                                                                ionisation, as therapeutic supplement in patients having specific deficiency in hombre. Production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hombre. may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1, in cohromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                  N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence.
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Pred. No. 2.7e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 2 A; 5 C; 7 G; 3 T; 0 other;
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Best Local Similarity 87.5%;
Marches 14; Conservative
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(GLAX )
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Shannon ME;

Chen W,

Hanzel DK, Rank DR,

Ji Y, Penn SG,

Gu Y,

(AEOM-) AEOMICA INC.

30-JAN-2001; 05-FEB-2001;

WPI; 2002-179446/23.

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1

Disclosure; SEQ ID 10730; 214pp; English.

o

Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber

Claim 4; Page 67; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, tumour angiogenesis, diabetic retinopathy, macular degeneration, cumour neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca vulgaris, angiofibroma of tubercous sclerosis, port-wine stains, Sturge Weber syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting comprises the use of one or more theraples to tumour conjunction with one or more of the treatment. Leukaemia or tumour conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of office reducing ERG activity in a cell, by contacting (I) is useful for reducing ERG activity in a cell, by contacting (I) with ENG, in the presence of a divalent confice and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically targeting genes that share homology with ERG gene or ERG fusion genes. ABK17354-ABK22719 represent nucleic acid, including antisense and centared port he invention.

Sequence 17 BP; 2 A; 0 C; 5 G; 10 U; 0 other;

Gaps 0; Length 17; Indels 0.9%; Score 12.8; DB 1; 31.2%; Pred. No. 2.7e+02; ative 9; Mismatches 2; 9; 5; Conservative Query Match Best Local Similarity Matches

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1325 TIGIAGAICTIGIGIT 1340 .:|:|:||: .:|:|
UUGUGGAUUUUGUGUU 16

à В ABK18357

RESULT 556

ABK18357 standard; RNA; 17 BP.

ABK18357;

09-APR-2002 (first entry)

Human ERG hammerhead ribozyme target sequence, Seq ID No 1004.

Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antidiarthritic; antiporiatic; virucid; osteopathic; vulnerary; cancer; lymphome; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leuxaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme; amberzyme.

Homo sapiens.

WO200188124-A2.

22-NOV-2001.

16-MAY-2001; 2001WO-US15866.

16-MAY-2000; 2000US-0572021. 

(RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.

Randi AM; Mclaughlin F, McSwiggen JA, Von Carlowitz I, Jarvis T,

WPI; 2002-082995/11.

Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome 

Claim 4; Page 77; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates corporations selected from cancer, lymphoma, Ewing's sarcoma, melanoma, conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, tumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic selecosis, port-whis stains, Sturge Weber syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. Leukaemia or tumour cumber conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or cannot such as Mg2+. (I) is useful for reducing ERG activity in a coll seases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect trapeting genes that share homology with ERG gene or ERG fusion genes. ABK1734-ABK2719 represent nucleic acids, including antisense and enzymetic nucleic acid molecules which regulate expression of ERG, and a cell of PCF related PCF related to the inventions which regulate expression of ERG, and a cell of ERG which a cell of the invention and mutations with regulate expression of ERG, and as cells or code enzymetic nucleic acids, including antisense and enzymetic nucleic acide which regulate expression of ERG, and related PCR primers of the invention.

Sequence 17 BP; 5 A; 6 C; 3 G; 3 U; 0 other;

. 0 0.9%; Score 12.8; DB 1; Length 17; 5.0%; Pred. No. 2.7e+02; ve 2; Mismatches 2; Indels 75.0%; Matches 12; Conservative Local Similarity Query Match

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713 CIGIGGCCCAGCAGCA 728 2 cuguedcccauchach 17 à g

ABK18359

ABK18359 standard; RNA; 17 BP

ABK18359; 

(first entry) 09-APR-2002 Human ERG hammerhead ribozyme target sequence, Seq ID No 1006.

Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Enving's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome; leukaemia; osteoporosis; DNAzyme; inozyme; amberzyme.

Homo sapiens

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the invention fraction to a nucleic acid molecule (1) is useful for treating conditions selected from cancer, lymphoma; Ewing's sarcoma, melanoma, conditions selected from cancer, lymphoma; Ewing's sarcoma, melanoma, cumour angiogenesis; diabetic retinopathy, macular degeneration, neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge Weber syndrome, Naipel-Tremaunay-Weber syndrome, Osle-Weber-rendu syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (1) under conditions suitable for the treatment. The method comprises the use of one or more therapies to the treatment. Leukaemia or tumour conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting (I) with RNA, in the presence of a divalent cation such as Mg2+. (I) is useful for diagnosis of conditions and diseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically the presence of ERG RNA in a cell. (I) is useful for specifically cargeting genes that share homology with ERG gene or ERG fusion genes. ABKI7354-ABK22719 represent nucleic acide, including antisense and enzymatic nucleic acid molecules which regulate expression of ERG, and acides and cenzymatic nucleic acid molecules which regulate expression of ERG, and acides and cenzymatic nucleic acide molecules which regulate expression of ERG, and acides and cenzymatic nucleic acide molecules which regulate expression of ERG, and acides and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a nucleic acid molecule (I) which down regulates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel polynucleotide which down regulates expression of Ets-related
                                                                                                                                                                                                                                                                                                                                       Von Carlowitz I, McSwiggen JA, Mclaughlin F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; Page 77; 149pp; English.
                                                                                                                                     16-MAY-2001; 2001WO-US15866.
                                                                                                                                                                                              16-MAY-2000; 2000US-0572021.
                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                              (GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-082995/11.
                       WO200188124-A2
                                                                                22-NOV-2001
                                                                                                                                                                                                                                                                                                                                       Jarvis T,
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Sequence 17 BP; 5 A; 5 C; 4 G; 3 U; 0 other;

Gaps ; 1) 13. Score 12.8; DB 1; Length 17; larity 75.0%; Pred. No. 2.7e+02; Conservative 2; Mismatches 2; Indels Query Match Best Local Similarity Matches 12; Conserv

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ABK18805 standard; RNA; 17 BP. RESULT 558 ABK18805,

(first entry) 09-APR-2002 ABK18805;

Human ERG DNAzyme target sequence Seq ID No 1452.

Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; 

tumour angiogenesis, diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme; amberzyme.

Homo sapiens.

WO200188124-A2.

22-NOV-2001.

Randi AM;

16-MAY-2001; 2001WO-US15866.

6-MAY-2000; 2000US-0572021. 

(RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.

Randi AM; Mclaughlin F, Von Carlowitz I, McSwiggen JA, Jarvis T,

WPI; 2002-082995/11.

Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome

Claim 4; Page 92; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, malanoma, tumour anglogenesis, diabetic retinopathy, macular degeneration, tumour anglogenesis, diabetic retinopathy, macular degeneration, neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca vulgaris, anglofibroma of tuberous sclerosis, port-whose stains, Sturge Weber syndrome, Ishkemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies the treatment in the new of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting the cell with (I). (I) is useful for cleaving RNA of ERG gene, by contacting (I) with RNA, in the presence of a divalent cation such as M24. (I) is useful for diagnosis of conditions and diseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically targeting genes that share homology with ERG gene or ERG fusion genes. ARX.7354-ABX22719 represent nucleic acids, including antisense and enzymatic nucleic acids, including antisense and enzymatic nucleic acids, including antisense and enzymatic nucleic acids, including antisense and related PCR primers of the invention.

Sequence 17 BP; 5 A; 7 C; 2 G; 3 U; 0 other;

Gaps 0; Length 17; Indels Query Match 0.9%; Score 12.8; DB 1; Best Local Similarity 87.5%; Pred. No. 2.7e+02; 0; Mismatches 14; Conservative Matches

; 0

TGATGCAGCTGGAGTT 17 셤

824 TGATGCAGCTGAAGCT 839

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ABK19286 standard; RNA; 17 BP ABK19286

RESULT 559

ABK19286;

09-APR-2002 (first entry)

Human ERG Amberzyme target sequence Seq ID No 1933.

Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous solerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme; amberzyme.

Homo sapiens.

WO200188124-A2.

22-NOV-2001

16-MAY-2001; 2001WO-US15866.

16-MAY-2000; 2000US-0572021

(RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.

Jarvis T, Von Carlowitz I, McSwiggen JA, Mclaughlin F, Randi AM;

WPI; 2002-082995/11.

Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome

Claim 4; Page 124; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Buning's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, theorems are accoma, myopic degeneration, arthritis, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge weber syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies to under conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting the cell with (I). (I) is useful for cleaving RNA of ERG gene, by contacting (I) with RNA, in the presence of a divalent cation such as Mg2+. (I) is useful for diagnostic tool to cation such as Mg2+. (I) is useful for diagnostic tool to cation such as Mg2+. (I) is useful for diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically cargeting genes that share homology with ERG gene or ERG funion genes.

ABK1754-ABK27194 represent nucleic acids, including and cargeting genes that share homology with ERG gene or ERG funion genes. enzymatic nucleic acid molecules which regulate expression of ERG, and related PCR primers of the invention.

Sequence 17 BP; 8 A; 5 C; 3 G; 1 U; 0 other;

Gaps . 0 Ouery Match 0.9%; Score 12.8; DB 1; Length 17; Best Local Similarity 81.2%; Pred. No. 2.7e+02; Indels 2; 1; Mismatches 13; Conservative Matches

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2 ACCCACAGAAGAUGAA 17

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RESULT 560 ABK19333

ABK19333 standard; RNA; 17 BP.

ABK19333;

(first entry) 09-APR-2002 Human ERG Amberzyme target sequence Seg ID No 1980.

Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme; amberzyme.

Homo sapiens

WO200188124-A2.

22-NOV-2001.

16-MAY-2001; 2001WO-US15866.

16-MAY-2000; 2000US-0572021.

(RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LID.

Von Carlowitz I, McSwiggen JA, Mclaughlin F, Randi AM; Jarvis T, 

WPI; 2002-082995/11.

Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome

Claim 4; Page 126; 149pp; English.

cumoun auglogenesis, diabetic retinopathy, macular degeneration, cumoun auglogenesis, mappic degeneration, arthritis, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge Weber syndrome, leukaemia, osteoporosis and wound healing. (1) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (1) under conditions suitable for the treatment. The method comprises the use of one or more therapies under conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treated by administering (1) to the patient in conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or coll, by contacting the cell with (1). (1) is useful for cleaving RNA of ERG gene, by contacting (1) with RNA, in the presence of a divalent of issue as more of ERG RNA, in the presence of a divalent of disabses related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (1) is useful for specifically caracting genes that share homology with ERG gene or ERG fusion genes. ABK17354-ABK2219 represent nucleic acids, including antisense and craymatic nucleic acid molecules which regulate expression of ERG, and craymatic nucleic acid molecules which regulate expression of ERG, and The invention relates to a nucleic acid molecule (I) which down regulates expression of an Bts-related gene [REG]. (I) is useful for treating conditions selected from cancer, lymphoma, Bwing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration,

Length 17; Indels 0.9%; Score 12.8; DB 1; ilarity 87.5%; Pred. No. 2.7e+02; Conservative 0; Mismatches 2: Query Match Best Local Similarity 14; Matches

A; 1 C; 7 G; 3 U; 0 other;

Sequence 17 BP; 6

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Gaps

. 0

RESULT 561

ABK25511 standard; DNA; 17 BP ABK25511, 

ABK25511;

(first entry) 09-APR-2002

Male-sterile plant producing genome altering oligonucleotide #411.

Chromosomal genomic alteration; genome altering oligonucleotide; PCR, ss, o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin-B; abiotic stress tolerance; improved nutritional value; hygromycin: primary amino acid over production; herbicide resistance; glyphosate resistance; imidazolinone herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; raiafane resistance; modified oil production; modified starch production; waxy starch; antered floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced plant; albino mutant; increased stearate production; albino plant; increased stearate production; albino plant; photosynthetic process.

Zea mays

Synthefic.

WO200192512-A2

06-DEC-2001

01-JUN-2001; 2001WO-US17672

01-JUN-2000; 2000US-208538P. 30-OCT-2000; 2000US-244989P. 27-MAR-2001; 2001US-0818875.

(UYDE ) UNIV DELAWARE

Rice MC, Kmiec EB, Gamper HB,

Kim J;

WPI; 2002-106307/14.

New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production

Claim 7; Page 94; 220pp; English.

The invention relates to an oligonucleotide for targeted alteration of a genetic sequence, which comprises a single-stranded oligonucleotide having a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications or single of o-methyl modification, an LNA modification, two or more phosphorothioate linkages on a terminus, or a combination of any two or more of these modifications. The oligonucleotides are useful for the oligonucleotides are useful for disconting repair or alteration of plant genetic information. The oligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abiotic stress tolerance, improved nutritional value (e.g. altering amino acid content of plants or conferring amino acid over production), herbicide resistance (e.g.

resistance, porphyric herbicide resistance or triazine resistance), disease resistance, modified oil production, modified starch production modified starch production of ways starch, altered floral morphology (e.g. male-sterile plants) or modified fatty acid content (e.g. reduced palmitate, increased stearate or reduced linolenic acid). The oligonuclecitides are also useful for producing albino mutants for the analysis of photosynthetic processes. This sequence represents a genome altering oligonuclecitide of the invention. and sulphonylurea herbicide glyphosate resistance, imidazolinone 

Sequence 17 BP; 5 A; 3 C; 5 G; 4 T; 0 other;

Gaps 0 Length 17; Indels 0.9%; Score 12.8; DB 1; 37.5%; Pred. No. 2.7e+02; ve. 0; Mismatches 2; Local Similarity 87.5%; les 14; Conservative Query Match Best Loca Matches

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ò 셤 RESULT 562 ABK2551

BP. ABK25512 standard; DNA; 17

ABK25512;

(first entry) 09-APR-2002 Male-sterile plant producing genome altering oligonucleotide #412.

Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss; o-methyl modification; LNA modification; phosphorothioate linkage; but registry lnA alteration; environmental tolerance; hygromychin-B; abiotic stress tolerance; improved nutritional value; hygromychin-B; amino acid over production; herbicide resistance; glyphosate resistance; imidazolinone herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; sulphonylurea herbicide resistance; modified oil production; modified starch production; waxy starch; altered floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced plant; albino mutant; increased stearate production; reduced linolenic acid production; photosynthetic process.

Zea mays. Synthetic.

WO200192512-A2

06-DEC-2001

01-JUN-2001; 2001WO-US17672.

01-JUN-2000; 2000US-208538P. 30-OCT-2000; 2000US-244989P. 27-MAR-2001; 2001US-0818875.

(UYDE ) UNIV DELAWARE.

Kim J; Gamper HB, Rice MC, Kmiec EB,

WPI; 2002-106307/14.

New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or oil production modified

Claim 7; Page 94; 220pp; English.

๙ οĘ The invention relates to an oligonucleotide for targeted alteration genetic sequence, which comprises a single-stranded oligonucleotide having a DNA domain. The DNA domain has at least one mismatch with

8X98999999999999999

chemical modifications of the oligonucleotide. The chemical modifications chemical modification, an LNA modification, two or more consist of onethyl modification, an LNA modification, two or more phosphorothicate linkages on a terminus, or a combination of any two or more of these modifications. The oligonucleotides are useful for directing repair or alteration of plant genetic information. The oligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abilotic stress tolerance, improved nutritional value (e.g. altering amino acid content of plants or conferring amino acid over production, herbicide resistance (e.g. glyphosate resistance, imidazolinone and sulphomylurea herbicide conferring amino acid outlined to triazine resistance, indazolinone and sulphomylurea herbicide resistance, conferring amino acid only production, modified starch production (e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatty acid content of e.g. reduced palmitate, increased stearate or reduced linolenic acid). The oligonucleotides are are also useful for producing albino mutants for the analysis of photosynthetic processes. This sequence represents a genome altering oligonucleotide of the invention.

Sequence 17 BP; 4 A; 5 C; 3 G; 5 T; 0 other;

Length 17; Query Match 0.9%; Score 12.8; DB 1; Length 1 Best Local Similarity 87.5%; Pred. No. 2.7e+02; Matches 14; Conservative 0; Mismatches 2; Indels 164 GATCCTCAAGGTCTCG 179 16 1 GATCCTCTAGATCTCG ð

Gaps

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ABK25527; ABK25527, 

ABK25527 standard; DNA; 17 BP.

09-APR-2002 (first entry)

Male-sterile plant producing genome altering oligonucleotide #427.

Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss; o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin-B; abiotic stress tolerance; improved nutritional value; hygromycin, primer; amino acid over production; herbicide resistance; glyphosate resistance; imidazolinone herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; triazine resistance; disease resistance; altered floral morphology; male-sterile production; waxy starch; modified floral morphology; male-sterile plant; albino mutant; increased stearate production; reduced plant; albino mutant; increased stearate production; reduced linolenic acid production; plant; increased stearate production; reduced linolenic acid production; photosynthetic process.

Oryza sativa.

Synthetic.

WO200192512-A2.

06-DEC-2001.

01-JUN-2000; 2000US-208538P. 30-OCT-2000; 2000US-244989P. 27-MAR-2001; 2001US-0818875.

01-JUN-2001; 2001WO-US17672.

(UYDE ) UNIV DELAWARE.

Kim J; Rice MC, Gamper HB, Kmiec EB,

WPI; 2002-106307/14.

New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production -Claim 7; Page 94; 220pp; English

The invention relates to an oligomucleotide for targeted alteration of a genetic sequence, which comprises a single-stranded oligomucleotide of having a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligomucleotide. The chemical modifications of the oligomucleotide. The chemical modifications of the oligomucleotides are useful for more of these modifications. The oligomucleotides are useful for directing repair or alteration of plant genetic information. The oligomucleotides are particularly useful for creating plants with desired of phenotypes, e.g. environmental or abilotic stress tolerance, improved nutritional value (e.g. altering amino acid content of plants or contenting amino acid over production, herbicide resistance, indazolinone and sulphomylurea herbicide (e.g. altering amino acid content of plants or glyphosate resistance, indazolinone and sulphomylurea herbicide (e.g. increased starch or production, modified starch production (e.g. increased starch or production of waxy starch), altered floral corpusional plants or modified fatty acid content of e.g. increased starch or production of waxy starch), altered floral corpusional plants or modified fatty acid content of e.g. increased starch or production of waxy starch), altered floral corpusionally should plants or reduced linolatic soft increased stearate or reduced linolatic for the analysis of photosynthetic processes. This sequence represents a genome of altering oligonucleotide of the invention. 

Sequence 17 BP; 5 A; 3 C; 5 G; 4 T; 0 other;

° Gaps ö 0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; ve 0; Mismatches 2; Indels Local Similarity 87.5%; 14; Conservative Query Match Matches

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ABK25528 standard; DNA; 17 BP. RESULT 564 **ABK25528** 

(first entry)

09-APR-2002

Male-sterile plant producing genome altering oligonucleotide #428.

Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss; o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair: DNA alteration; environmental tolerance; hygromycin; priabiotic stress tolerance; improved mutritional value; hygromycin; primer; amino acid over production; herbicide resistance; glyphosate resistance; imidazolinone herbicide resistance; sulphonylurea herbicide resistance; modified oil production; modified starch production; modified starch production; waxy starch; altered floral morphology; male sterile plant; albino mutant; modified fatty acid content; reduced palmitate production; albino plant; modified fatty acid content; reduced palmitate production; albino plant; increased stearate production; reduced linolenic acid production; photosynthetic process

Oryza sativa, Synthetic WO200192512-A2.

36-DEC-2001,

01-JUN-2001; 2001WO-US17672.

genotypes

Nishida M;

Moriya S,

Kmiec EB,

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The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (BAEJ0512-ABEJ31809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of hudividuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, dances, Langerhams islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when
                                                                                                                                                                                   Human leukocyte antigen (HLA) typing, useful for judging HLA of individuals e.g. by determining immunogenetic differences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, human leukocyte antigen, HLA; genotype; polymorphism;
immunogenetic; transplantation; genetic disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human HLA genotyping oligonucleotide SEQ ID NO 1053.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%; Score 12.8; DB 1;
87.5%; Pred. No. 2.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matsumura Y,
                                                                                         Matsumura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 1 A; 7 C; 6 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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                                                                                                                                                                                                                                                                                       Claim 10; Page 199; 345pp; Japanese.
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                                                                                         Ichihara T,
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transplanting between them -
                                                                                                                                                                                                             of individuals e.g. by detern
transplanting between them -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.
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                 (NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.
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nes 14; Conservative
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                                                                                         Kagiya T,
                                                                                                                                          WPI; 2002-122074/16
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                                                                                         Inoko H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 566
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             radying a DNA domain. The DNA domain has at least one distinction with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications of the oligonucleotide. The chemical modifications of phosphorothicate linkages on a terminus, or a combination of any two or more of these modifications. The oligonucleotides are useful for a directing repair or alteration of plant genetic information. The cligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abiotic stress tolerance, improved nutritional value (e.g. alterning amino acid content of plants or conferring amino acid over production), herbicide resistance (e.g. glyphosate resistance, imidazolinone and sulphonylurea herbicide (e.g. resistance, modified oil production), modified starch production (e.g. increased starch or production, modified starch production (e.g. increased starch or production, modified fatty acid content (e.g. reduced palmitate, increased stearete or reduced linolenic acid). The oligonucleotides are also useful for producing albino mutants for the allowally so of plants of the sequence represents a genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an oligonuclectide for targeted alteration of genetic sequence, which comprises a single-stranded oligonuclectide having a DNA domain. The DNA domain has at least one mismatch with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                         New oligonucleotides with modified nuclease-resistant termini, usefu
for creating plants with desired phenotypes, e.g. stress tolerance,
improved nutritional value, herbicide or disease resistance, or
modified oil production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human HLA genotyping oligonucleotide SEQ ID NO 562.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         altering oligonucleotide of the invention.
                                                                                                                                          Kim J;
                                                                                                                                                                                                                                                                                                                                                                 Claim 7; Page 94; 220pp; English
                                                                                                                                             Rice MC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 164 GATCCTCAAGGTCTCG 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL31073 standard; DNA; 17 BP
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01-JUN-2000; 2000US-208538P.
30-OCT-2000; 2000US-244989P.
                                              27-MAR-2001; 2001US-0818875
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14; Conservative
                                                                                         (UYDE ) UNIV DELAWARE
                                                                                                                                             Gamper HB,
                                                                                                                                                                                           WPI; 2002-106307/14.
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Gaps

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Indels

Length 17;

Nishida

Moriya S,

21-MAR-2002

ABL31073;

RESULT 565

ABL31073

Query Match

Matches

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06-DEC-2001

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The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (BAB130512ABE131809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, dangerham sislet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals. Gaps 0; 0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; tive 0; Mismatches 2; Indels Sequence 17 BP; 4 A; 4 C; 7 G; 2 T; 0 other; Ouery Match
Best Local Similarity 87.5%
Local 14; Conservative

531 GGAGCAGCTGGGTGCC 546 geaechecreachecce 16 g à

Human HLA genotyping oligonucleotide SEQ ID NO 1144. ABL31655 standard; DNA; 17 BP (first entry) 21-MAR-2002 ABL31655; RESULT 567 ABL31655 

Human; human leukocyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.

Homo sapiens

06-DEC-2001.

01-JUN-2001; 2001WO-JP04662,

01-JUN-2000; 2000JP-0164798.

(NISN ) NISSHINBO IND INC.

(SYST-) SYSTEM RES INC.

Inoko H, Kagiya T, Ichihara T,

Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them -WPI; 2002-122074/16.

Matsumura Y, Moriya S, Nishida

Claim 10; Page 310; 345pp; Japanese.

The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligomuclectides (AB15012-AB151809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals.

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Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them -
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                                          Gaps
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                                                                                                                                                                                                         Human, human leukocyte antigen, HLA; genotype; polymorphism;
immunogenetic; transplantation; genetic disease; ss.
                        Length 17;
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                                           Indele
                                                                                                                                                                                       Human HLA genotyping oligonucleotide SEQ ID NO 1160.
                        0.9%; Score 12.8; DB 1;
87.5%; Pred. No. 2.7e+02;
Live 0; Mismatches 2;
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     Sequence 17 BP; 5 A; 8 C; 3 G; 1 T; 0 other;
                                          0;
                                                              CCACCCGGACGCCGAC 210
                                                                                2 ccaccaddacdccrac 17
                                                                                                                                                                                                                                                                                                01-JUN-2001; 2001WO-JP04662.
                                                                                                                                                                                                                                                                                                                  01-JUN-2000; 2000JP-0164798.
                                                                                                                                ABL31671 standard; DNA; 17
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                                                                                                                                                                      (first entry)
                                 Local Similarity 87.5
les 14; Conservative
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                                                                                                                                                                                                                                         Homo sapiens.
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                                                                195
                                                                                                                                                     ABL31671;
                          Query Match
                                                                                                             RESULT 568
                                            Matches
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The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (ABL30512-ABL31809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals. Sequence 17 BP; 4 A; 4 C; 7 G; 2 T; 0 other;

Claim 10; Page 313; 345pp; Japanese.

0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; ive 0; Mismatches 2; Indels 531 GGAGCAGCTGGGTGCC 546 ilarity 87.5%; Conservative Similarity 14; Query Match Local Matches

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Gaps

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1 GGAGCAGCTGAGAGCC 16

g à

Renault

Mengaud J,

06-DEC-2001

Inoko H,

21-MAR-2002

ABL31720;

us09904568-3.rng

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The present invention relates to eps operons from Streptococcus thermophilus (see ABA01431-ABA01448). Proteins encoded by the eps operon function in exopolysaccharide (BPS) synthesis. The operons are useful for producing chimeric eps operons, for optimising production of EPS in lactic acid bacteria. EPS impart texture, mouth feel and rheological properties to fermenced food products (e.g. yoghurt). They function as thickeners, to provide free-flowing and creamy texture, and may also have biological activities beneficial to health. The present sequence is a PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                                                                                                                                                                                                                                                                                                                                                   New nucleic acid fragments containing exopolysaccharide operon, useful e.g. for increasing exopolysacchride synthesis in lactic acid bacteria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer, which was used in an example from the present invention
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Pred. No. 2.7e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                      Fremaux C,
                                                                                                                                                                                  (INRG ) INRA INST NAT RECH AGRONOMIQUE. (DANO-) CIE DANONE SA GERVAIS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 31; 144pp; French
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                                                                                                                                                                                                                                                                      Rallu F, Besancon-Yoshpe I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-SEP-2002; 2002WO-IB04208.
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                                                                                                                                             18-APR-2000; 2000FR-0004972.
                                                                                                   18-APR-2001; 2001WO-FR01199
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Best Local Similarity 87.5.
Best Local 14; Conservative
                                                                                                                                                                                                                            (RHOD ) RHODIA CHIM.
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                 WO200179500-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them -
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immunogenetic, transplantation; genetic disease; ss.
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                                                                                                                                               Human HLA genotyping oligonucleotide SEQ ID NO 1209.
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                        ABL31720 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.
                                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kagiya T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-122074/16.
                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer; ss.
                                                                                                                                                                                                                                                                                              WO200192572-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Streptococcus
                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-FEB-2002
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Streptococcus thermophilus

ABA01457

570

ABA01457 RESULT

Query Match

à g

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Gaps

. 0

Length 17; Indels

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Sequence 17 BF; 2 A; 3 C; 1 G; 11 T; 0 other;
WPI; 2003-313353/30.
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the interaction retailed to a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them under highly stringent conditions, or the complement of any of the invention are useful as probes and primers isolated nucleic acids of the invention are useful as probes and primers of detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the nucleic acids, cells containing the nucleic and so polypeptides directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of these therapy. This polymeticide sequences of the invention can be used in gene therapy. This polymuclecide sequence represents a tumour suppression related human fukutin oligonuclectide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel isolated 17 mer nucleic acid sequence,
New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells
                                                                                                                                                                                                                                                                                         Disclosure; Page 203; 720pp; French
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Gaps
                                ..
0
0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; ive 0; Mismatches 2; Indels
                                                                1139 ATGCCTTTTTTTT 1154
                                                                                           Arccrititititi 17
                                   Conservative
                Best_Local Similarity
Matches 14; Conserv
                                                                                              N
      Query Match
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Tumour suppression related human fukutin oligo SEQ ID No 1486. BP. ABT35849 standard; DNA; 17 (first entry) 12-JUN-2003 ABT35849; RESULT 572 ABT35849/ 

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; senee; tumour; cell degeneration; cancer; Alzheimer's disease; ecizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.

WO2003025175-A2. Homo sapiens,

L7-SEP-2002; 2002WO-IB04208.

27-MAR-2003

MOLE-) MOLECULAR ENGINES LAB

.7-SEP-2001; 2001FR-0011978

Tuijnder M; Amson R, relerman A,

WPI; 2003-313353/30.

New isolated nucleic acid, useful for treating viral diseas associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells

Disclosure; Page 206; 720pp; French

The invention relates to a novel isolated 17 mer nucleic acid sequence, or given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after consecutive nucleotides from the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents, cond for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the polypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for prevention and/or treatment of viral ciseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell containing the expression of the 17 mer nucleic acids in containing the expression of the 17 mer nucleic acids in gene contained sequences of the invention can be used in gene contained by the polypeptides can also be used to generate antibodies, and chips. The nucleic acid sequences of the invention can be used in gene contained by the polypeptide sequence represents a tumour suppression contained therapy. This polymucleotide of the invention. 

Sequence 17 BP; 4 A; 6 C; 4 G; 3 T; 0 other;

Gaps . o Length 17; 2; Indels 0.9%; Score 12.8; DB 1; 87.5%; Pred. No. 2.7e+02; 0; Mismatches Local Similarity 87.5 les 14; Conservative Query Match Matches

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Dp

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RESULT 573

ABT36006 standard; DNA; 17 BP. (first entry) 12-JUN-2003 ABT36006; ABT36006

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzhaimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds. Tumour suppression related human fukutin oligo SEQ ID No 1643.

WO2003025175-A2. Homo sapiens 

17-SEP-2002; 2002WO-IB04208

27-MAR-2003.

17-SEP-2001; 2001FR-0011978.

(MOLE-) MOLECULAR ENGINES LAB.

Tuijnder M; Amson R, relerman A,

WPI; 2003-313353/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related

Thu Jan

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The invention relates to a movel isolated 1, met nuclet data sequence, the invention relates to a movel isolated 1, met nuclet data sequence, optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated mucleic acids of the invention are useful as probes and primers or detecting, identifying quantifying and/or amplifying a nucleic acid, e.g. as one component of recombinant polypeptides. Any of the nucleic acid, onlypeptides, compound the containing the nucleic acids, polypeptides, and of recombinant polypeptides. Any of the nucleic acids, polypeptides, octor or antibodies directed against the polypeptides are useful for production of pharmaceuticials for prevention and/or treatment of viral diseases that are characterised by development of tumours or call degeneration, specifically cancer but also Alzheimer's disease and diseases that are characterised by development of tumours or call degeneration, specifically cancer but also Alzheimer's disease and cation propression of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chieses. The nucleic acid sequence represents a tumour suppression related human fukutin oligonucleotide of the invention.
                                                                                                                          The invention relates to a novel isolated 17 mer nucleic acid sequence,
polypeptides, antibodies and transfected cells
                                                                Disclosure; Page 225; 720pp; French.
      õ
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ô Gaps ; 0 Query Match

0.9%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 6 A; 2 C; 5 G; 4 T; 0 other;

ABT37187 standard; DNA; 17 ABT37187; RESULT 574 ABT37187 

BP.

(first entry) 12-JUN-2003

Tumour suppression related human fukutin oligo SEQ ID No 2824.

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.

Homo sapiens

WO2003025175-A2.

27-MAR-2003

17-SEP-2002; 2002WO-IB04208.

(MOLE-) MOLECULAR ENGINES LAB.

17-SEP-2001; 2001FR-0011978.

Tuljnder M; Telerman A, Amson R,

WPI; 2003-313353/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells

The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding NNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the optypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral degeneration, specifically cancer but also Alzehemer's diseases and degeneration, specifically cancer but also Alzehemer's diseases and cancer but also Alzehemer's diseases and parties of these and parties are useful and parties an both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention. Disclosure, Page 363; 720pp; French. 

Sequence 17 BP; 8 A; 2 C; 5 G; 2 T; 0 other;

Gaps .; 0 Length 17; Indels 0.9%; Score 12.8; DB 1; 37.5%; Pred. No. 2.7e+02; ve 0; Mismatches 2; 87.5%; Local Similarity 87.5 les 14; Conservative Query Match Best Loca Matches

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à d

ABT37634 standard; DNA; 17 BP 12-JUN-2003 (first entry) ABT37634; RESULT 575 ABT37634 

Tumour suppression related human fukutin oligo SEQ ID No 3271.

Cytostatic, virucide, neuroprotective, nootropic, neuroleptic, gene chip, antieense, sense, tumour, cell degeneration, cancer, Alzheimer's disease, schizophrenia, protein chip, gene therapy, tumour suppression, human fukutin, ds.

Homo sapiens.

WO2003025175-A2

27-MAR-2003.

17-SEP-2002; 2002WO-IB04208.

17-SEP-2001; 2001FR-0011978.

Tuijnder M; relerman A, Amson R,

(MOLE-) MOLECULAR ENGINES LAB.

WPI; 2003-313353/30.

New isolated mucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells

Disclosure; Page 416; 720pp; French

the invalidation related to a large and the containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers to detecting, identifying, quantifying and/or amplifying a nucleic acid, or detecting, identifying, quantifying and/or amplifying a nucleic acid, or one component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, colls containing the nucleic acids, colls containing the polypeptides are useful for proparation of pharmaceuticials for prevention and/or treatment of viral degeneration, specifically cancer but also Alzheimer's disease and degeneration, specifically cancer but also Alzheimer's disease and string amples is useful for diagnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polymericatide sequence repersents a tumour suppression contained therefore the proposed the invention of protein contained by the procession of the invention. invention relates to a novel isolated 17 mer nucleic acid sequence, 8888888888888888888888888888

Sequence 17 BP; 2 A; 7 C; 4 G; 4 T; 0 other;

Gaps .. Query Match

0.9%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

164 GATCCTCAAGGTCTCG 179 GATCCCCACGGTCTCG 16

g à

Tumour suppression related human fukutin oligo SEQ ID No 3442. ABT37805 standard; DNA; 17 BP (first entry) 12-JUN-2003 ABT37805; RESULT 576 

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.

Homo sapiens

WO2003025175-A2.

27-MAR-2003

17-SEP-2001; 2001FR-0011978. 17-SEP-2002; 2002WO-IB04208.

(MOLE-) MOLECULAR ENGINES LAB.

Tuijnder M; Telerman A, Amson R,

WPI; 2003-313353/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells -

Disclosure; Page 436; 720pp; French.

The invention relates to a novel isolated  $17\ \mathrm{mer}$  nucleic acid sequence, given in the specification, a sequence containing at least 15

consecutive nucleotides from the 17 mer sequence, a sequence, a sequence, at least 80 % identity to the 17 mer sequence, a sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acid, polypeptides, vectors containing the nucleic acids, cells containing the polypeptides in useful for preparation of pharmaceuticals for prevention and/or treatment of viral degeneration, specifically cancer but also Alzheimer's disease and schill seases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schill schild and antibodies can also be used to generate antibodies, and both the polypeptides can also be used to generate antibodies, and chips. The nucleic acid sequence represents a tumour suppression cherapy. This polynucleotide sequence represents a tumour suppression crelated human fukutin oligonucleotide of the invention. 

Sequence 17 BP; 6 A; 2 C; 4 G; 5 T; 0 other;

ö Gaps . 0 Match 0.9%; Score 12.8; DB 1; Length 17; Local Similarity 87.5%; Pred. No. 2.7e+02; Loss 14; Conservative 0; Mismatches 2; Indels Query Match Matches

983 CAGTCCCATTCAGATC 998 16 cagirccarraagarc 1 g à

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ABT38097 standard; DNA; 17 RESULT 577 ABT38097 

BP.

ABT38097;

(first entry) 12-JUN-2003 Tumour suppression related human fukutin oligo SEQ ID No 3734.

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antiesnse; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.

Homo sapiens

WO2003025175-A2.

27-MAR-2003.

17-SEP-2002; 2002WO-IB04208.

17-SEP-2001; 2001FR-0011978

(MOLE-) MOLECULAR ENGINES LAB.

Tuijnder M;

Amson R,

Telerman A,

WPI; 2003-313353/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related associated with tumors and cell degeneration, a polypeptides, antibodies and transfected cells

Disclosure; Page 470; 720pp; French.

The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a

sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, as one component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the propagation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell diseases. The polypeptides can also be used to generate antibodies, and patient samples is useful for diagnosis and/or prognosis of these contains the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention. 

Sequence 17 BP; 2 A; 5 C; 6 G; 4 T; 0 other;

ch 0.9%; Score 12.8; DB 1; Length 17; 1 Similarity 87.5%; Pred. No. 2.7e+02; 14; Conservative 0; Mismatches 2; Indels 1066 CCCATCAGGCAGGCTC 1081 CCCATCAGGGAGGATC 1 Local Similarity 16 Query Match Best Loca Matches ð 엄

ABT39257 standard; DNA; 17 ABT39257; 

B.

(first entry) 12-JUN-2003

Tumour suppression related human fukutin oligo SEQ ID No 4894.

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.

Homo sapiens

WO2003025175-A2.

27-MAR-2003

17-SEP-2002; 2002WO-IB04208.

17-SEP-2001; 2001FR-0011978.

(MOLE-) MOLECULAR ENGINES LAB

Tuijnder M; Amson R, Telerman A,

WPI; 2003-313353/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells

Disclosure; Page 606; 720pp; French

The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 sidentity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel

isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (atti)sense reagences, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for disgnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and chips. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention. 

Sequence 17 BP; 7 A; 3 C; 3 G; 4 T; 0 other;

Gaps . 0 Length 17; Query Match

0.9%; Score 12.8; DB 1; Length 1

Best Local Similarity 87.5%; Pred. No. 2.7e+02;

Matches 14; Conservative 0; Mismatches 2; Indels 14; Conservative

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16 ATACTGCTTTGAGATC 855 ATACCGCTTTGAGGTC

à g

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Gaps .. RESULT 5

ABT39551 standard; DNA; 17 BP.

ABT39551;

12-JUN-2003 (first entry)

Tumour suppression related human fukutin oligo SEQ ID No 5188.

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.

Homo sapiens.

WO2003025175-A2.

27-MAR-2003.

17-SEP-2002; 2002WO-IB04208.

17-SEP-2001; 2001FR-0011978.

(MOLE-) MOLECULAR ENGINES LAB.

Tuijnder M; Telerman A, Amson R,

WPI; 2003-313353/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells -

Disclosure; Page 640; 720pp; French.

given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, The invention relates to a novel isolated 17 mer nucleic acid sequence,

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e.g. as one component of a gene chip, in vitro as (anti)sense reagents, polypeptides. Any of the nucleic acids, polypeptides, wectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for disquosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention.
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Sequence 17 BP; 2 A; 3 C; 1 G; 11 T; 0 other;

.; 0 Gaps ; 0.9%; Score 12.8; DB 1; Length 17; 17.5%; Pred. No. 2.7e+02; Indels 0; Mismatches 87.5%; Matches 14; Conservative Best Local Similarity Query Match

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580

ABT39720 standard; DNA; 17 BP.

ABT39720

(first entry) 12-JUN-2003 Tumour suppression related human fukutin oligo SEQ ID No 5357.

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds. fukutin;

Homo sapiens.

WO2003025175-A2.

27-MAR-2003

17-SEP-2002; 2002WO-IB04208.

17-SEP-2001; 2001FR-0011978.

(MOLE-) MOLECULAR ENGINES LAB.

Tuijnder

Amson R,

Telerman A,

WPI; 2003-313353/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells

Disclosure; Page 660; 720pp; French.

The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid e.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, 

polypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these disease. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention. 8¥866666666668

Sequence 17 BP; 4 A; 2 C; 5 G; 6 T; 0 other;

Gaps . 0 Length 17; Score 12.8; DB 1; Length 1 Pred. No. 2.7e+02; 0; Mismatches 2; Indels 0; 0.9%; Local Similarity 87.5 nes 14; Conservative Query Match Matches

·: 0

 $\delta$ 셤 RESULT 581

ACA06264 standard; RNA; 17 BP. ACA06264

ACA06264;

03-JUN-2003 (first entry)

NFKB sub-unit modulating inozyme substrate #83.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; oesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; oervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; cyclophosphanide; docetaxel; docetaxel; cisplatin; methotrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatorid arthritis; restenosis; Crohn's disease; obesity; iscanamia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; HANDER STANDER STANDER

Homo sapiens.

US2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-0864785.

94US-0291932. 92US-0987132. 94US-0245466. 96US-0777916 15-AUG-1994; 07~DEC-1992; 18-MAY-1994; 23-DEC-1996;

₽. STINCHCOMB D MCSWIGGEN J. (STIN/) (MCSW/)

DRAPER K G.

(DRAP/)

Draper KG; Mcswiggen J, DT, Stinchcomb

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases

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regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRS), where (I) is an inozyme, zinzyme, G-leaver or ambezzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the leavel of REL-A. (I) is useful for cleaving RNA condition associated with the leavel of REL-A. contisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oespecially Mg^2+. The enzymatic and muscate, colorectal, brain, oespecially scomach, bladder, pancreatic, prostate, colorectal, brain, oespecial, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, gencitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as
                                                                                         invention describes an enzymatic nucleic acid molecule (I) which down
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocarchial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 3 A; 4 C; 8 G; 2 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      enzymatic nucleic acid molecule.
                          Claim 3; Page 28; 72pp; English
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0; Gaps 0.9%; Score 12.8; DB 1; Length 17; 31.2%; Pred. No. 2.7e+02; ve 1; Mismatches 2; Indels ilarity 81.2%; Pr Conservative 1; 479 AGGACTGCCGAGACGG 494 1 AGGACUGCCGGGAUGG 16 Local Similarity hes 13; Conserv Query Match Best Loca Matches g

ACA06326 standard; RNA; 17 BP. (first entry) 03-JUN-2003 ACA06326; RESULT 582 ACA06326 

NFKB sub-unit modulating inozyme substrate #145.

G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; brostate cancer; colorectal cancer; brostate cancer; colorectal cancer; brades cancer; pancreatic cervical cancer; stomach cancer; bladder cancer; pancreatic cancer; lymphoma; plioma; miltidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; gemcitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection;

Homo sapiens.

US2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-0864785.

94US-0291932 15-AUG-1994;

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The invention describes an engineer and miscours of muchest factor kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The engyment on coled acid molecule is adapted to treat concern and is useful for down-regulating KBL-A activity in a cell, for treating a patient having a condition associated with the level of RBL-A. (I) is useful for clearing RNA comprising a sequence of RBL-A gene, in the presence of a divalent cation, especially MG-2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stromach, bladder, parcreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug theraphs such as monoclonal antibodies, RBL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, gemcitabine or radiation therapy. The enzymatic and antisense nucleic caid molecules are also useful for treating inflammatory disease such as recalls of the seaso, lungus, multiple solerosis, transplant/graft celection, gene therapy applications, ischaemia/reperfusion injury central nervous system (CNS) and myocardial), glomerulomephilits, session multiple is allowed disease or section methods are interested to the seaso of the section multiple solerosis, transplant/graft central nervous system (CNS) and myocardial), glomerulomephilits, session multiple and seaso or sepsion allowed disease or section multiple may inflammatory be a section of a section multiple and section multiple and section and may see a section of section and myocardial), glomerulomephilits, session multiple may not section inflammatory because or section methods and may be and section multiple may not section inflammatory because or section may not section the section multiple may not section inflammatory of a section of a section of the section section in the section of a section of section section in the section of section section sect
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 0 A; 11 C; 3 G; 3 U; 0 other;
                                                                                                                                                                                                                                               Draper KG;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 3; Page 29; 72pp; English.
                                                                                                                                                                                                                                               Stinchcomb DT, Mcswiggen J,
                   94US-0245466.
                                                                                                                    (STIN/) STINCHCOMB D T. (MCSW/) MCSWIGGEN J.
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                                                                                                                                                                                DRAPER K G.
                       18-MAY-1994;
23-DEC-1996;
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Gaps .; 0 Length 17; 0.9%; Score 12.8; DB 1; Length 1 5.0%; Pred. No. 2.7e+02; ve 2; Mismatches 2; Indels Local Similarity 75.0%; nes 12; Conservative Query Match **Matches** 

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ò g ACA06585 standard; RNA; 17 BP 03-JUN-2003 (first entry) ACA06585;

NFKB sub-unit modulating inozyme substrate #404.

G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; cerophageal cancer; stomach cancer; cancer; pancreatic cancer; cervical cancer; pancreatic cancer; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemocherapy; paclitexel; doceaxel; cipplatin; methofrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; gemcitabine; radiation therapy; inflammatory disease; asthma; diabetes; Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; 

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rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia;
                                                            gene therapy, autoimmune disease; lupus, multiple sclerosis, sepsis, transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection;
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Homo sapiens.

US2002177568-A1.

28-NOV-2002

2001US-0864785. 23-MAY-2001; 94US-0291932. 92US-0987132. 94US-0245466. 96US-0777916. 18-MAY-1994; 23-DEC-1996; L5-AUG-1994; 07-DEC-1992

STINCHCOMB D STIN/)

MCSWIGGEN J. MCSW/) MCSWIGGEN J DRAP/) DRAPER K G. Draper KG; Stinchcomb DT, Mcswiggen J,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases

Claim 3; Page 33; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor regulates expression of a sequence encoding a subunit of nuclear factor regulates B (NRKB), where (I) is an inozyme, d-deaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a paintent having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or untidung resistant cancer. The method involves use of other drug charapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, displatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, genicitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as chematod arthritis, restenosis, resting, cobesity, autoimmune disease, lupus, multiple solerois, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury centerion, gene therapy applications, ischaemia/reperfusion injury sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel 

Sequence 17 BP; 3 A; 6 C; 6 G; 2 U; 0 other;

Gaps ; 0 Query Match, 0.9%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 2.7e+02; Matches 14; Conservative 0; Mismatches 2; Indels Query Match

· 0

570 GCTCCAGCAGGCCCTC 585 N 17 GCTGCAGCAGGGCCTC

RESULT 584 ACA06587

ACA06587 standard; RNA; 17 BP. EX

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-Cleaver; amberzyme; cancer; REL-A acitvity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; oesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methorexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephitis; allergic airway inflammation; inflammatory bowel disease; infection; NFKB sub-unit modulating inozyme substrate #406. 94US-0291932. 92US-0987132. 94US-0245466. 23-MAY-2001; 2001US-0864785. 96US-0777916. 03-JUN-2003 (first entry) ₽. STIN/) STINCHCOMB D MCSW/) MCSWIGGEN J. US2002177568-A1. Homo sapiens. 15-AUG-1994; 23-DEC-1996; 07-DEC-1992; 18-MAY-1994; 28-NOV-2002. ACA06587; 

Draper KG; Stinchcomb DT, Mcswiggen J,

DRAP/) DRAPER K G.

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases

Claim 3; Page 33; 72pp; English.

regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRB), where (I) is an inozyme, cinzyme, d-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG-2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, crivial, head and neck, ovarian cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxal, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, catarexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, come therapy applications, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (NR) and myocardial), glomerulomephritis, espesies are also useful for the substrate of a novel encymatic nucleic acid molecule. The invention describes an enzymatic nucleic acid molecule (I) which down

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention describes an enzymatic nucleic acid molecule (I) which down
                                                                                                                                                                                                                                                                                                                                                                                                     G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; human; oesophageal cancer; cancer; colorectal cancer; brain cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sespais; transplant/graft rejection; reperfusion injury; glomerulonephritis; reperfusion injury; glomerulonephritis; allergic airway inflammatory bowel disease; infection;
                                                                                                                                                                                                                                                                                                                                                                   Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;
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                                                                                Gaps
                                                                              0;
                                    ch 0.9%; Score 12.8; DB 1; Length 17; Similarity 68.8%; Pred. No. 2.7e+02; 11; Conservative 3; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                   NFKB sub-unit modulating inozyme substrate #472.
 Sequence 17 BP; 2 A; 6 C; 5 G; 4 U; 0 other;
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                                                                                                               837
                                                                                                                                                                                                                                        ACA06653 standard; RNA; 17 BP
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92US-0987132.
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                                                                                                             822 CCTGATGCAGCTGAAG
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С
                                    Query Match
Best Local Similarity
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07-DEC-1992;
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multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, gencitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restenosis, asthma. Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; cesophageal cancer; stomach cancer; bland and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxal; docetaxel; cisplatin; methotrexate; chemotherapy; paclitaxal; docetaxel; cisplatin; methotrexate; cyclophosphanide; docetaxel; cisplatin; daterexate; cyclophosphanide; docetaxel; cisplatin; daterexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; pheumatorid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; repertusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection;
   glioma or
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 head and neck, ovarian cancer, melanoma, lymphoma,
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92US-0987132.
94US-0245466.
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                                                                                                                                                                                                                                                                                                                                                                                                                              17 GGAGCAGGAGCTGGGT
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07-DEC-1992;
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23-DEC-1996;
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The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inozyme, d-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A, enc. (I) is useful for cleaving RNA comprising a sequence of REL-A, enc. (I) is useful for cleaving RNA comprising a sequence of REL-A, enc. (I) is useful for cleaving RNA comprising because of REL-A, enc. (I) is useful for cleaving RNA comprising because of REL-A, enc. (I) the presence of a divalent cation, especially MG^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lumg, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug cervical, as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, are also useful for treating inflammatory disease such as cid molecules are also useful for treating inflammatory disease, obseity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischemia/reperfusion injury confirmation. This sequence reperfusion, inflammatory bowel disease or inflection. This sequence reperfusion, inflammatory bowel disease or inflation. This sequence reperfusion, inflammatory bowel disease or inflation. This sequence is the substrate of a novel of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases enzymatic nucleic acid molecule. Claim 3; Page 36; 72pp; English 

Sequence 17 BP; 4 A; 7 C; 3 G; 3 U; 0 other;

Gaps 0; Query Match 0.9%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 2.7e+02; Matches 14; Conservative 0; Mismatches 2; Indels

0;

335 CTGGTGATAGTCACAG 350 Credretrasscaes 2 17 ద

ACA06858 standard; RNA; 17 BP ACA06858; RESULT 587 ACA06858 

03-JUN-2003 (first entry)

NFKB sub-unit modulating inozyme substrate #677.

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G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; breast cancer; human; oesophageal cancer; stomach cancer; colorectal cancer; brain cancer; cervical cancer; stomach cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemocherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphamide; doxorubhi; fluorouracil carboplatin; edatrexate; recursing arthritis; restenosis; Crohn's disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;

sapiens ОШОН

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;

NFKB sub-unit modulating inozyme substrate #678.

G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; oesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer;

cervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate;

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-0864785.

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regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The arzymatic nucleic acid molecule is adapted to treat configuration. The arzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, ossophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxal, docetaxal, cisplain, methorizate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide are also useful for treating inflammatory disease such as cidential arthritis, restenosis, asthma, Cronn's disease, diabetes, obesity, autoimmune disease, lupus, multiple solerosis, transplant/graft controlled to the controlled or rediations, isohamial/reperfusion injury
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Best Local Similarity 87.5%; Pred. No. 2.7e+02;
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                 94US-0291932.
92US-0987132.
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96US-0777916.
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                                                                                                                                                                  (DRAP/) DRAPER K G.
                                                                                                                                                                                                      Stinchcomb DT,
               15-AUG-1994;
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The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor regulates expression of a sequence encoding a subunit of nuclear factor configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprisaing a sequence of REL-A gene, in the presence of a divalent cation, especially MG^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, colorectal, as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, genicitabline or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as colosis, aschmanial/reperfusion injury (central nervous system (CNS) and mycoradial), glomerulonephritis, respense to inflammatory bowel disease or infection. This sequence represents the substrate of a novel
               gemcitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion inlury; glomerulomephritis; allergic airway inflammation; inflammatory bowel disease; infection;
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doxorubin, fluorouracil carboplatin; edatrexate;
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92US-0987132.
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96US-0777916.
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DRAP/) DRAPER K G.
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07-DEC-1992;
18-MAY-1994;
23-DEC-1996;
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Gaps
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                                  0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; ive 0; Mismatches 2; Indels
Sequence 17 BP; 1 A; 9 C; 1 G; 6 U; 0 other;
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937 GAGAAGAGGTGTGAGC 952
                                          16 GAGAAGAGGGGAGAGC 1
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ACA07770/c RESULT 589

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; cervical cancer; prostate and neck cancer; bladder cancer; pancreatic cancer; cervical cancer; plomach and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methorrexare; chemotherapy; paclitaxel; docetaxel; cisplatin; deterexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatorid arthritis; restenosis; Crohn's disease; obstiv; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; NFKB sub-unit modulating zinzyme substrate #169. ACA07770 standard; RNA; 17 BP. 94US-0291932. 92US-0987132. 94US-0245466. 96US-0777916. Stinchcomb DT, Mcswiggen J, 23-MAY-2001; 2001US-0864785 (first entry) STINCHCOMB D T. MCSWIGGEN J. DRAPER K G. US2002177568-A1. Homo sapiens. 03-JUN-2003 15-AUG-1994; 07-DEC-1992; 18-MAY-1994; 23-DEC-1996; 28-NOV-2002. ACA07770; (/NILS) (DRAP/) MCSW/) 

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases

Draper KG;

WPI; 2003-340953/32.

Claim 3; Page 40; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor regulates expression of a sequence encoding a subunit of nuclear factor configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, head and neck, ovarian cancer, melanoma, lymphoma, glioma or cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-Aspecific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, captures or chemmatoid arthritis, restenosis, asthma, crohm's disease such as consity, autoimmune disease, lupus, multiple sclerosis, transplant/graft crejection, gene therapy application, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, cespesis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel

enzymatic nucleic acid molecule.

antisense nucleic acid molecules are useful for treating breast, lung,

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The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2^+. The enzymatic and
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                                                                          Length 17;
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                                                                      0.9%; Score 12.8; DB 1;
ilarity 87.5%; Pred. No. 2.7e+02;
Conservative 0; Mismatches 2;
                                     Sequence 17 BP; 2 A; 6 C; 6 G; 3 U; 0 other;
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                                                                                                                                              570 GCTCCAGCAGGCCCTC 585
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                                                                      Query Match
Best Local Similarity
Matches 14; Conserv
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07-DEC-1992;
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prostate, colorectal, brain, despendent, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glloma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edertexate, gencitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such a rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, septence represents the substrate of a novel enzymatic nucleic acid molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a screening method for selecting compounds and their salts inhibiting the expression of core binding factor alpha subunit 1 (CBFA1) type II, by measuring the amount of mRNA encoding CBFA1 type II exon 0 present in cells expressing CBFA1, in the presence and absence of the test compound. CBFA1 has cytostatic activity. The method can be used for the treatment, prevention and diagnosis of cancer (including cancer of the colon, breast, lung, prostate, oesophagus, liver, gall-bladder, spleen, kidney, pancreas, uterus, testis, thyroid, bladder or brain, and leukaemia) and inhibition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Screening compounds inhibiting expression of core binding factor alpha subunit 1 type II for prevention and treatment of cancer and metastasis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   factor alpha subunit 1; CBFA1; cancer; metastasis;
                                                                                                                                                                                                                                                                                                                                                           Length 17;
                                                                                                                                                                                                                                                                                                                                                                                               2; Indels
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                                                                                                                                                                                                                                                                                                                                                       Score 12.8; DB 1;
Pred. No. 2.7e+02;
2; Mismatches 2;
                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 3 A; 8 C; 4 G; 2 U; 0 other;
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leukaemia; PCR primer; ss.
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Best Local Similarity 75.0
Matches 12; Conservative
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17 AGCTGAGATGGCGCCA
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of cancer proliferation and metastasis. The present sequence represents a PCR primer for human CBFA1 type II, which is used in an example from the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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                                                                                                              Length 17;
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0.9%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
                                                                    Sequence 17 BP; 4 A; 8 C; 3 G; 2 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                         Human K-Ras DNAzyme substrate #684.
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06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
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                                                                                                                                                                                                                                                                                                          ABZ60572 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                 679 GIGGIATITGGGAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-140484/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200297114-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J;
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ABZ60572;

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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HRR2. K-Ras, H-Ras, N-Ra human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-Heumatic activity. The nucleic acid molecules are useful for reducing HRR2. K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ65216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66520 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
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                                                                                                                                                                                                                                                             Human, ribozyme, short interfering RNA, siRNA, HER2, K-Ras, enzymatic nucleic acid, H-Ras, N-Ras, HIV, cytostatic, anti-HIV, anti-rheumatic; cancer, AIDS, ss.
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                                                                                                                                                                                                  Human K-Ras DNAzyme substrate #1283.
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06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
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ABZ61171 standard; RNA; 17
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                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-MAR-2003 (first entry)
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Matches 14; Conserv
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ABZ61760/c
ID ABZ617
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acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule encoding HER2, K-Ras, H-Ras, M-Ras, and in-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ25889 - ABZ66516, ABZ66531, ABZ66530 - ABZ66516, ABZ66531, and AIDS. sequences for the human ribozymes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel short interfering RNA (siRNA) nucleic
                                                                                                                                                                                                                                                                                                          Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
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enzymatic nucleic acid, H-Ras, N-Ras, HIV, cytostatic, anti-HIV,
anti-rheumatic, cancer, AIDS, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 12.8; DB 1;
Pred. No. 2.7e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                   Claim 58; Page 121; 185pp; English.
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06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
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                                                                                    29-MAY-2001; 2001US-294140P.
06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
                                       29-MAY-2002; 2002WO-US16840
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                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC.
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Best Local Similarity 87.5.
Frankling 14; Conservative
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                                                                                                                                                                                                                         Mcswiggen J;
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05-DEC-2002
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enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
                                                          Human, ribozyme, short interfering RNA, siRNA; HER2; K-Ras;
enzymatic nucleic acid, H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 58; Page 121; 185pp; English.
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                  Human H-Ras DNAzyme target #551.
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10-SEP-2001; 2001US-318471P.
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Best Local Similarity 87.5
Matches 14; Conservative
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RESULT 595 ABZ61761

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Gaps

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Length 17; 2; Indels

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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HRR2, K-Ras, H-Ras, N-Rahuman immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-Heumatic activity. The nucleic acid molecules are useful for reducing HRR2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ5989 - ABZ65816, ABZ65531, ABZ6520 - ABZ65850 - ABZ6585 represent substrate/target sequences for the human ribozymes of the invention.
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                                               Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
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                                                                                                                    Claim 4; Page 139; 185pp; English
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06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
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                  WPI; 2003-140484/13
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Best Local Similarity
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expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic canner, and HIV infection, and AIDS. The sequences shown in ABZ53989 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66585 represent substrate/target sequences for the human ribozymes of the invention.
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                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fibroblast growth factor; hybridisation; laser procedures; wascular smooth muscle cell; proliferation; SMC; vascular stenosis; post a mojoplasty restenosis; atherosclerosis; cardiac hypertrophy; organ transplant; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New anti-sense polynucleotide(s) to fibroblast growth factor receptor - used for inhibiting vascular smooth muscle cell proliferation, partic. for treating restenosis
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                                                                                                                                                                                                                                          Ouery Match
0.9%; Score 12.8; DB 1;
Best Local Similarity 81.2%; Pred. No. 2.7e+02;
Matches 13; Conservative 1; Mismatches 2;
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                                                                                                                                                                                                           Sequence 17 BP; 4 A; 2 C; 8 G; 3 U; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense oligonucleotide for mouse FGF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dixon RAF, Rege AA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3; Page 9; 53pp; English.
                                                                                                                                                                                                                                                                                                                    475 GGGGAGGACTGCCGAG 490
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ70348 standard; DNA; 18
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-DEC-1992;
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15-FEB-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ70348;
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AAT28333;
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AAT28333/
                       SON COURSE OF CO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer 1 and 2 (AAQ86978-79) were used to amplify a target nucleotide sequence from methicillin resistant S. aureus (MRSA). The target is detected with a probe specifically derived from the mech gene of S. aureus and S. epidermidis. The mech gene product has no homology with known PBPs (penicillin-binding proteins). The new probes allow for the rapid identification of all MRS, eradicating need for labour intensive in vitro cultivation and physiological assays.
                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                             MRSA; methicillin resistant Staphylococcus aureus; probe;
hybridisation; mecA; MRSE; Staphylococcus epidermis; primer; PCR;
polymerase chain reaction; ss.
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                                       0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.9%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 2.9e+02; Live 0; Mismatches 2; Indels
                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rabbit CETP hairpin ribozyme target sequence #191.
Pred. No. 2.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 7 A; 1 C; 4 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                         Primer 1 to amplify MRSA target DNA.
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                                                                                                                 CCCCACATCCCAGTTC 17
                  87.5%;
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Best Local Similarity 87.5
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Springer
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Staphylococcus aureus
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                       Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DE4338119-A1,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-NOV-1993;
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                                     Matches
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AAT50699-T50754 represent target sequences for the rabbit cholesterol ester transfer protein (CETP) hairpin ribozymes (see AAT50643-T50698).

CETP is a 74 m Glycoprotein that facilitates neutral lipid transfer to be between plasma lipoproteins. The numbering of the targets refers to binds to 4-6 mucleotides 5', and a variable number 3' of this site. The ribozymes are able to cleave mRNA from the gene encoding CETP, thereby blocking synthesis and/or expression of the mRNA. By inhibiting CETP, the reverse cholesterol transport (RCT) pathway can be inhibited (or bliminated) thereby preventing the reduction in size density of the high density lipoproteins (HDL), prolonging HDL half life, and therefore increasing HDL levels. The ribozymes can be used to treat conditions consisted with abnormal levels of CETP, specifically atherosclerosis; peripheral vascular disease, hyporbetalipoproteinaemia, vascular complications of diabetes, transplant, atherectomy and angioplastic restenosis. By inhibiting CETP, the levels of HDL and low density lipoproteins (LDL), and the HDL:LDL ratio are favourably altered (a decrease in LDL levels, and a corresponding increase in HDL levels). The ribozymes can also be used diagnostically to study genetic drift and material and in diseased cells, and to detect CETP mRNA, As the ribozymes can also be used diagnostically to study genetic drift and material and material and the ribozymes can also be used diagnostically to study genetic drift and material and the ribozymes can also be used diagnostically and the ribozymes can also be used diagnostically the ribozymes for the ribozymes can also be used diagnostically and the ribozymes can also be used diagnostically to study genetic drift and material and the ribozymes can also be used diagnostically and the ribozymes can also be used diagnostically the ribozymes and the ribozymes can also be used diagnostically and the ribozymes can also ribozymes can also be used diagnostically and the ribozymes can also ribozymes can also be used diagnosticall
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neutral lipid transfer; plasma lipoprotein; atherosclerosis; atherectomy; reverse cholesterol transport; high density lipoprotein; therapy; CBTP; familial hypercholesterolaemia; dyslipidaemia; hypoalphalipoproteinaemia; peripheral vascular disease; hyperbetalipoproteinaemia; RCT; inhibitor; angioplastic restenosis; low density lipoprotein; diabetes; HDL; rabbit;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mutations in diseased cells, and to detect CETP mRNA. As the ribozymes target specific regions of the CETP gene, they have low non-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New ribozyme(s) for cleaving cholesterol ester transfer protein mRNA - useful for preventing or treating initial development, progression or regression of vascular diseases, esp. familial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
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87.5%; Pred. No. 2.9e+02;
tive 0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (WARN ) WARNER LAMBERT
                                                                                                                                                                                                                                                                                                                                                                 Oryctolagus cuniculus
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Synthetic

us09904568-3.rng

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Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Primer ES11 (AAT30032) is based on the conserved amino acids CEGCKA(G) of the DNA binding domain of a typical nuclear receptor. It was used with antisense primer ES15 (AAT30034) for the PCR amplification of human colMa prepd. from osteosarcoma SAGS-2/B10 cells. A second round of PCR utilised primers ES12 (AAT30033) and ES15. A product was obtd. that was used to screen the osteosarcoma ibitarry, resulting in isolation of a clone (AAT30031) coding for a novel steroid hormone receptor, NER (AAR98140).
                                                                                                                               NEX receptor; potentiator; steroid hormone receptor; G-protein coupled receptor; nerve growth factor; Alzheimer disease; ocular hypertension; schilzophrenia; distonia; tardive dyskinesia; Gilles de la Tourette syndrome; polymerase chain reaction; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         steroid hormone receptor NER activators - for potentiating
                                                                                                                                                                                                                                                                                                                                               /*tag= a
/note= 'sequence differs from ES11 sequence on
pages 32 and 52 (see also AAT30035)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%; Score 12.8; DB 1; Length 18;
77.8%; Pred. No. 2.9e+02;
Lve 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse flt-1 VEGF receptor hairpin ribozyme substrate #87.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                activity of modulator of G-protein coupled receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schmidt A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 3 A; 3 C; 7 G; 3 T; 2 other;
                                                                                                                             potentiator; steroid hormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MEDI-) MEDICAL COLLEGE PENNSYLVANIA.
(MERI ) MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rodan GA,
                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 31; 63pp; English.
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                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Holloway MK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                             *tag=
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                                                                                  NER PCR primer ES11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                       misc difference
                                                                                                                             NER receptor;
                                          19-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-0CT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                WO9613257-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Friedman E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-JUL-1999
                                                                                                                                                                                                                primer; ss.
                                                                                                                                                                                                                                                          Synthetic
  AAT30032;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX75628;
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ID AAX7

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              are oligonucleotides of the invention. These sequences can be modified to become more nuclease resistant, using phosphorothicate, problem modified links. To screen for modified mustile sequences that inhibit cell proliferation, cultured smooth mustile cells that are arrested in the GO phase, are induced to proliferate in the presence of the multi-G sequence. The cultured smooth muscle cells used in this method are attraction medium, followed by transfer to a normal growth arrest is achieved on a starvation medium, followed by transfer to a normal growth medium to induce proliferation. The compounds that provide over 50% inhibition at a set dosage are selected as being useful for inhibiting vascular restencis. The multi-G oligonucleotides are used to inhibit proliferation of smooth muscle cells, such as to prevent
                                                   Multi-G oligonucleotide; antisense sequence; c-myb; nuclease resistant; phosphorothioate linkage; phosphorodithioate linkage; inhibitor; therapy; cell proliferation; smooth muscle cell; proliferation protein; vascular restenosis; arterial restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           arterial restenosis. These sequences are not antisense sequences, but are thought to work in a similar way. The sequences are thought to act by binding to proteins involved in the proliferation process. Compounds containing these multi-G oligonucleotides are not toxic, and their effect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT28317-T283347 represent multi-G oligonuclectides. AAT28332-T28335 are multi-G oligonuclectides with inosine substitutions. These sequences are oligonuclectides of the invention. These sequences can be modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New modified oligo:nucleotide(s) contg. consecutive guanine residues - inhibit proliferation of smooth muscle cells, esp. to prevent arterial restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12.8; DB 1; Length 18;
Pred. No. 2.9e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 0 A; 5 C; 7 G; 4 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cell proliferation is fully reversible.
             Multi-G oligonucleotide hu SCR (I2G2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fisher EF;
                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 46; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        886 CAGGAGCTGCGGTACAGC 903
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ID AAT30032 standard; DNA; 18 BP.
XX
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                                                                                                                                                                                                                                                                          mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Farrell CL,
                                                                                                                                                                                                                                                 *tag=
                                                                                                                                                                                                                                                                                                                *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1996-209848/21
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                                                                                                                                                                                                                         modified base
                                                                                                                                                                                                                                                                                       modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Burgess TL,
                                                                                                                                                                                                                                                                                                                                                                                                                       18-APR-1996.
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Vogel RL;

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Gaps

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expression

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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient freeferby human) having a condition associated with the level of the fms-like tyrosine kinase i (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Maize, corn, Zea mays, delta-9 desaturase, GBSS, target, substrate, granule bound starch synthase, hammerhead ribozyme; hairpin ribozyme; modulation; gene expression; transgenic plant; cleavage; canola plant; caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
                                                                                                                                                                                                      Nucleic acid molecule modulating VEGF receptor(s) gene expressimRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Granule bound starch synthase hairpin substrate SEQ ID NO:609.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ribozyme which modulates plant gene expression - preferably
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 12.8; DB 1;
Pred. No. 2.9e+02;
5; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           McSwiggen JA,
Zwick MG;
                                                                                                                        Stinchcomb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 2 A; 4 C; 6 G; 6 U; 0 other;
                                                                                                                        Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           O, Guo L,
Young SA,
                                                                                                                                                                                                                                                                                       Claim 4; Page 94; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1311 GTAGCCAGGTGCTTTT 1326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
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96US-0584040.
95US-0005974.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX62734 standard; RNA; 18
                                                                               (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                          McSwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Folkerts
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                                                                                                                                                                 WPI; 1997-259017/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1997-202224/18
                                                            (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DOWC ) DOWELANCO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-JUL-1999
11-JAN-1996;
26-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9710328-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-JUL-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Edington BE,
Merlo PAO, 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-MAR-1997.
                                                                                                                        Escobedo J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX62734;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       605
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VBGF). A patient (preferably human) having a condition associated with the level of the fins-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid archritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX6775 to AAX7575 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vascular endothelial growth factor receptor; VEGF receptor; flt-1; tDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%; Score 12.8; DB 1; Length 18; 56.2%; Pred. No. 2.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human flt1 VEGF receptor hairpin ribozyme substrate #62.
                                                                                                                                                                                                                                                                                                                                                  Stinchcomb D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 2 A; 5 C; 6 G; 5 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5; Mismatches
                                                                                                                                                                                                                                                                                                                                                  McSwiggen J, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; Page 188; 218pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1311 GTAGCCAGGTGCTTTT 1326
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                                                                                                                                                                                                                                                                                   (CHIR ) CHIRON CORP.
(RIBO-) RIBOZYME PHARM INC.
                     foetal liver kinase 1; ss.
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                                                                                                     WO9715662-A2
                                                                                                                                                                                    25-OCT-1996;
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                                                                                                                                                                                                                                                                                                                                                  Escobedo J,
                                                                                                                                         01-MAY-1997.
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Query Match

Matches

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RESULT 604

AAX70294

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Gaps

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Indels

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Merlo

(I) can be used

(preferably a maize or canola plant).

in a plant

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                                                                                                        with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDNA sequence encoding maize Delta-9 desaturase. (1) can be used to modulate expression of a gare, in a plant (preferably a maize or canola plant). (1) can be used to modulate expression of a gare, in a plant (preferably a maize or canola plant). (1) can be used to modulate caffeine synthesis in a coffee plant, nicotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, chrysanthemum or marigold plant or lignin production in a tobacco, separation or plant or pine plant or lignin production in a tobacco,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes an enzymatic nucleic acid molecule (I) with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDNA sequence encoding maize Delta-9 desaturase. (I) can be used to modulate expression of a gene, preferably Delta-9 desaturase or a granule bound starch synthase (GBSS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Maize; corn; Zea mays; delta-9 desaturase; GBSS; target; substrate; granule bound starch synthase; hammerhead ribozyme; hairpin ribozyme; modulation; gene expression; transgenic plant; cleavage; canola plant; caffeine synthesis; coffee plant; micotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
                                                                                             present invention describes an enzymatic nucleic acid molecule (I)
                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound starch
modulates expression of DELTA-9 desaturase or granule bound starch
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Granule bound starch synthage hairpin substrate SEQ ID NO:629.
                                                                                                                                                                                                                                                                                                                                              0.9%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 2.9e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Merlo DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                McSwiggen JA,
Zwick MG;
                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 4 A; 6 C; 5 G; 3 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 42; Page 84; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Folkerts O, Guo L, cokut TA, Young SA,
                                                       Claim 42; Page 84; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВР
                                                                                                                                                                                                                                                                                                                                                                                                                        884 TCCAGGAGCTGCGGTA 899
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                      synthase in maize or canola
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(RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                      14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1997-202224/18.
                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Edington BE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JUL-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Merlo PAO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX62754;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zea mays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 606
                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                           The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX62754
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A novel method has been developed for identifying an unknown allele of a polyalielic gene. The method involves: (a) contacting the unknown allele with a panel of probes, each of which recognises a sequence motif that is present in some alleles of the polyalielic gene but not in others; (b) observing which probes recognise the unknown allele so as to obtain a fingerprint of the unknown alleles. The present sequence represents a specifically claimed probe for use in the method where the polyalielic with fingerprints of known alleles. The present sequence represents a specifically claimed probe for use in the method where the polyalielic of comparing the nummalian MHC genes is gene is a human leukocyte antigen class I gene. The method can be used for gene is a human leukocyte antigen class I gene. The method can be used for genes such as mammalian MHC genes, in mammals, TAP, LMP, ras, conclassical HAA class I genes, human complement factor genes C4 and C2, and the HAA complex, and genes located in mitochondrial DNA, bacterial chromosomes and viral DNA. The method is particularly useful for matching the alleles of the HAA genes in a prospective donor and a prospective recipient in tissue or organ transplantations. The method can also be used in paternity testing, in forensic medicine, as a
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            to modulate caffeine synthesis in a coffee plant, nicctine production a tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a tobacco, aspen, poplar or pine plant.
                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ĕ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying unknown allele(s) of a poly:allelic gene using panel of probes each recognising a sequence motif present in some allele(s) useful for donor matching in tissue transplantation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human leukocyte antigen; HLA; probe; tissue transplantation; MRG gene; major histocompatibility complex; paternity test; forensic medicine; haematological malignancy; inherited disorder; adoptive immunotherapy; identification; ss.
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                                                                                                                                                                                0.9%; Score 12.8; DB 1; Length 18; 8.8%; Pred. No. 2.9e+02; ve 3; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human leukocyte antigen class I gene URSTO probe 531-548.
                                                                                                                                           Sequence 18 BP; 2 A; 7 C; 5 G; 4 U; 0 other;
                                                                                                                                                                                                     Pred. No. 2.9e
; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BONE MARROW TRUST ANTHONY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Madrigal A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 5; Page 19; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                               AAT94805 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                   220 CGAGCTCCTCAGCCTC 235
                                                                                                                                                                                                                                                                                                        17
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                                                                                                                                                                                                       68.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-310717/28.
                                                                                                                                                                                                     Best Local Similarity
Matches 11; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NOLA-) NOLAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-NOV-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9720197-A2
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                                                                                                                                                                                                                         11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT94805;
                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                      RESULT 607
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follow up technique in treatment of haematological malignancies or inherited disorders, in adoptive immunotherapy, and in identification of bacteria and viruses. The method can provide for the identification of alleles of the polyallelic genes using a limited number of selected recurring motif probes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A method for treating airway disease in a subject has been produced, which involves the topical administration of an essentially adenosine free antisense oligonucleotide (ON) to the airway epithelium of the subject. The present sequence is an antisense oligonucleotide the specific for the human endothalin. The method can be used to treat alrway diseases such as cystic fibrosis, asthma, chronic obstructive pulmonary disease, bronchitis and other airway diseases characterised by an inflammatory response. By eliminating adenosine from the antisense ON, its liberation upon antisense degradation is prevented, thereby preventing adenosine-induced bronchoconstriction in patients
                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treatment of airway diseases such as asthma - by topically applying adenosine-free antisense oligo:nucleotide to airway epithelium of
                                                                                                                                                                                   ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Asthma; airway epithelium; adenosine free; cystic fibrosis; chronic obstructive pulmonary disease; bronchitis; ss.
                                                                                                                                          Query Match
0.9%; Score 12.8; DB 1; Length 18;
Best Local Similarity 87.5%; Pred. No. 2.9e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human endothelin-1 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 0 A; 4 C; 3 G; 11 T; 0 other;
                                                                                                          Sequence 18 BP; 5 A; 4 C; 7 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 5; Page 38; 71pp; English.
                                                                                                                                                                                                                   546
                                                                                                                                                                                                                                                                                                                                             AAT76410 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                 1 GGAGCAGCTGAGAGCC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-0474497,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      with hyper-reactive airways.
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                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                 531 GGAGCAGCTGGGTGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nyce JW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-051871/05.
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                                                                                                                                                                                                                                                                                                                                                                                                                15-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-DEC-1996.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                               AAT76410;
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                                                                                                                                                                                                                                                                                                         RESULT 608
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AAX09121-X10268 are allele-specific oligonucleotide primers used in the isolation of various biallelic polymorphic markers found in the human genome (represented in AAX10269-X12937). These primers can be used in a method for determining polymorphic forms in an individual for use in e.g. forenaics, paternity testing or for phenotypic typing for diseases such as agammaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, typerropley, Wiskott-Aldrich syndrome, Pabry's disease, familial hypercotholesterolemia, polycystic kidney disease, hereditary haemorrhagic telangiectasia, familial colonic polyposis, Bilers-Danlos syndrome, osteogenesis imperfecta, acute intermittent porphyria, autoimmune diseases, inflammation, cancer, diseases of the nervous system, infection by pathogenic microorganisms, and characteristics such as longevity, appearance (e.g. baldness, obesity), strength, speed, andurance, fertility, and susceptibility or receptivity to particular drugs or therapeutic treatments. The isolated polymorphic nucleic acid segments can also be used to produce medicaments for the treatment or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                     detection; phenotypic typing; characteristic; infection; hereditary; autoimmune disease; cancer; inflammation; drug; therapy; medicament; treatment; marker; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated nucleic acid segments from the human genome - used for determining polymorphic forms for use in e.g. forensics, paternity testing or phenotypic typing for disease
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                                                                                                                                                      Polymorphism; biallelic; human; forensic; paternity testing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 18;
                                                                                                                   Human biallelic polymorphic marker downstream primer #495.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.9%; Score 12.8; DB 1;
77.5%; Pred. No. 2.9e+02;
ve 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 5 A; 6 C; 4 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                       (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 16; Page 212; 310pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1285 CATACAGTIGCTCAGC 1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wang D;
                BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      prophylaxis of such diseases.
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              AAX10189 standard; DNA; 18
                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-286974/25.
                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                 WO9820165-A2
                                                                                                                                                                                                                                                                                                                                                                   05-NOV-1997;
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                                                                                  24-MAR-1999
                                                                                                                                                                                                                                                                                                                                  14-MAY-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hudson T,
                                                                                                                                                                                                                                             Synthetic
                                               AAX10189;
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AAV95056
ID AAV9509
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AAX10189
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Gaps

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Length 18; Indels

0.9%; Score 12.8; DB 1; 87.5%; Pred. No. 2.9e+02;

Mismatches

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Conservative

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Matches

Query Match Best Local Similarity

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24-FEB-1999

AAV95056;

11-JUN-1998.

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AAV35388-V35414 are primers used in a method which results in a recombinant vaccinia virus comprising of a gag gene from a retrovirus such as HIV-1 or HIV-2, fused to a DNA fragment containing an epitope region (30-300 bases in length) of a retroviral gene other than the gag gene may be altered so as to produce a gag protein modified from the natural sequence by the addition, deletion or substitution of at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hypervariable region; ENV protein; vaccinia virus; gag gene; retrovirus; vaccines; infection; protection; primer; ss.
                                                                                                                                                                                                                                                            This is the nucleotide sequence of a PCR primer used in the method of the invention, involving the use of novel apoptosis-related DNAs and proteins. The inventions can be used as diagnostic reagents for apoptosis e.g. (monoclonal) antibodies for the protein, as a reagent in immunohistological staining, as apoptosis inhibitors. It can also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Recombinant vaccinia virus containing fusion HIB gag gene -
production in host cells of gag protein for use as vaccine
                                                                                                                                                                    Novel apoptosis-related DNAs and proteins - for diagnosis, preventing or treating diseases associated with apoptosis
                                                                                                                                                                                                                                                                                                                                                                                                                                    0.9%; Score 12.8; DB 1; Length 1
87.5%; Pred. No. 2.9e+02;
ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                             in immunohistological staining, as apoptosis inhibit
be used for treatment of apoptosis-related diseases.
                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 1 A; 0 C; 2 G; 15 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NINA-) JAPAN NAT INST INFECTIOUS DISEASES (JAPG ) NIPPON ZEON KK.
                                                                                                                                                                                                                           Example 1; Page 47; 70pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 64; 84pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HIV-1 gag protein DNA primer #4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1144 TITITITITITIGGA 1159
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                                                   (KYOW ) KYOWA HAKKO KOGYO KK
                   97JP-0050302
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kojima A, Kurata T,
                                                                                                                                WPI; 1998-495844/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-312481/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
               05-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV35391;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                        Sakaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the synthesis and/or expression of interleukin (II)-ZR gamma encoded RNA. AAV93889 to AAV94574 represent specifically claimed ribozymes, and AXV94575 to AAV93526 represent specifically claimed substrate sequences from the present invention. The ribozymes can be used for the treatment of, e.g. graft rejection, autoimmune disease, cancer, psoriasis, allergy and other inflammatory conditions. The ribozymes are also used to induce tolerance in a recipient to alloantigen from a donor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present sequence invention describes ribozymes targeted to modulate
                                                                                                                  Human; IL-2 receptor g-chain; interleukin 2 receptor gamma chain;
hammerhead ribozyme; hairpin ribozyme; substrate; expression; cancer;
autoimmune disease; psoriasis; allergy; inflammatory disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR; primer; amplification; apoptosis; antibody; inhibition; ss; immunohistological staining.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ribozymes targeted to interleukin 2 - useful for treating e.g. cancer, autoimmune disease and allergies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                               Mouse IL-2 receptor g-chain substrate position 399.
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11.2%; Pred. No. 2.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 3 A; 7 C; 4 G; 4 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; Page 44; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV54165 standard; cDNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         640
                                                                                                                                                                                                                                                                                                                                                                                                                                              Stinchcomb DT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18
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                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC
                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     625 GACCAGCTCCAGGAGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.9
Best Local Similarity 81.2
Matches 13; Conservative
                                                                                                                                                                               graft rejection; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-33332/29
                                                                                                                                                                                                                                                                                                                                                                                                                                              McSwiggen JA,
                                                                                                                                                                                                                                                       WO9824913-A2
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The .

for

21-DEC-1998

AAV54165

RESULT 611

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AAV54165

11-SEP-1998,

Synthetic

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Gaps

0;

Length 18;

least 1 amino acid residue. The fusion gene is inserted into a region of a vaccinia virus not essential to its propagation, to give a recombinant vaccinia virus vector which is used to transform a host cell (such as Hela, Vero, VEF, rabbit kidney RK13 or human myeloma TK-143 cells). Upon culturing the host cell produces particulate structures containing the pusion aga protein. The recombinant vaccinia virus or the fusion gag protein and be used in the production of vaccines for protecting against infection with retroviruses such as HIV. 88888888888888

Sequence 18 BP; 1 A; 1 C; 1 G; 15 T; 0 other;

Gaps .; 0 Query Match

0.9%; Score 12.8; DB 1; Length 18;
Best Local Similarity 87.5%; Pred. No. 2.9e+02;
Matches 14; Conservative 0; Mismatches 2; Indels 1141 GCCTTTTTTTTTTT 1156 Gérrérérérerent 17

RESULT 613 AAZ33132/c

AAZ33132 standard; DNA; 18 BP.

AAZ33132;

26-JAN-2000 (first entry)

Treponema pallidum Msp variable region sense PCR primer S8.

Treponema pallidum; syphilis; vaccine; treponemal disease; Msp; major sheath protein; bejal; gingivitis; periodontal disease; pinta; yaws; PCR primer; ss.

Synthetic.

Preponema pallidum

WO9953099-A1

21-OCT-1999.

99WO-US07886. 09-APR-1999;

98US-0058968 10-APR-1998;

(UNIW ) UNIV WASHINGTON

Cameron CES Van Voorhis WC, Lukehart SA, Centurion-Lara GA,

WPI; 1999-620445/53.

proteins useful in vaccines against syphilis and other treponemal

Claim 3; Page 86; 200pp; English.

diseases

Novel

genes encode a gircerophosphodiester phosphodiesterase (Gpd), and proteins with homology to major outer sheath D15/oma87 homologue, and proteins with homology to major outer sheath (Msp) proteins of i denticola. Also described are: (1) an isolated protein capable of inducing a protective immunologic response to in an effective amount to an animal host; (2) a method of identifying a T. p. pallidum vaccine candidate; (3) a method of inducing a protective immune response against T. pallidum; (4) a method for analysing a sample of DNA to determine whether it originated from analysing a sample of DNA to determine whether it originated from analysing a sample of DNA to determine whether a first and a second clinical isolate of T. p. pallidum are the same or different. The clinical isolate of T. p. pallidum are the same or different. The clinical isolate of T. p. pallidum are the same or different. The clinical diseases, e.g. syphilis, bejal, pinta, yaws, gingivitis, and periodontal diseases. The methods may be used to identify vaccine other Treponemal diseases. The methods may be used to identify vaccine The present invention describes novel Treponema pallidum genes. These 

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candidates, and to determine the origin of a treponemal nucleic acid. AAZ33104 to AAZ33156, and AAY52773 to AAY52831, represent sequences used in the exemplification of the present invention.
                                                                                                                      Gaps
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                                                                                        Length 18;
                                                                                                                    Indels
                                                                                       0.9%; Score 12.8; DB 1;
87.5%; Pred. No. 2.9e+02;
                                                          Sequence 18 BP; 2 A; 8 C; 6 G; 2 T; 0 other;
                                                                                                                      Mismatches
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                                                                                                                                                523
                                                                                                                                                                             2
                                                                                                                                                508 GGGGTCAGCGCCAACC
                                                                                                                                                                           17 dedercadecercadec
                                                                                                                    14; Conservative
                                                                                    Query Match
Best Local Similarity
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                                                                                                                    Matches
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Cellular inhibitor of apoptosis-2 phosphorothioate antisense oligo #27. Identification; genetic target; gene modulation; human; probe; antisense oligonucleotide; phosphorothioate; PCR primer; nucleotide sequence-based technology; antisense drug discovery; (first entry) target validation; ss 26-JAN-2000 AAZ41035; 

BP.

AAZ41035 standard; DNA; 18

AAZ41035

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Homo sapiens. Synthetic

WO9953101-A1

21-OCT-1999.

99WO-US08268 13-APR-1999; 98US-0081483. 13-APR-1998; 28-APR-1998;

(ISIS-) ISIS PHARM INC.

Brooks DG; Sasmor HM, Freier SM, Vickers TA; Baker BF, McNeil J, Cowsert LM, band Wyatt JR,

WPI; 1999-620446/53.

Identifying compounds which modulate expression of nucleic acids, used to provide compounds having defined physical, chemical or bioactive properties, e.g. antisense activity

Example 21; Page 100; 264pp; English.

the expression of a target nucleic acid (tNA) sequence via binding of the expression of a target nucleic acid (tNA) sequence via binding of the expression of a target nucleic acid (tNA) sequence via binding of the compounds with the tNA sequence. The method comprises generating a library of virtual compounds in silico according to defined criteria. Also described are: (1) a method of tNA sequence via binding of the own with the tNA sequence comprising a tNA sequence via binding of the ONS with the tNA sequence comprising a tNA sequence via binding of the own with the tNA sequence comprising criteria, and evaluating in silico the binding of the virtual ONS with the tNA according to defined criteria; and (2) a method of defining a set of compounds that modulate the expression of a tNA sequence via binding of the compounds with the tNA. The methods can be used for the generation and identification of synthetic compounds having defined phinding of the compounds with the tNA. The methods can be used for the generation and identification of synthetic compounds having defined physical, chemical or bioactive properties. Information gathered from a ssays of such compounds is used to identify nucleic acid sequences that are tractable to a variety of nucleotide sequence-based technologies, e.g. antisense drug discovery and target validation. AAZ40852 to

55 X 8

RESULT 615 AAZ22129

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and proteins. When the CGICE gene is mutated it is responsible for best's macular dystrophy (BMD). Polynucleotides encoding CGICE are useful for diagnosing whether a patient carries a mutation in the CGICE gene. Normal and mutated CGICE proteins are useful for in order identifying activators and/or inhibitors of these proteins, in order to treat BMD. The CGICE gene offers a simpler and cheaper method of diagnosing BMD without the need for the presence of the patient. The gene may also be useful to discovering the genetic cause of age-related macular dystrophy. The present sequence represents a PCR primer for the human GCICE CDNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cute asthma, allergy, asthma, impeded respiration, respiratory distress syndrome; pain, cystic fibrosis, pulmonary hypertension; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes human and mouse CGICE polynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense oligonucleotide, multiple target, antisense treatment, impaired respiration, inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis;
                                                                                                                                            CGICE; Best's macular dystrophy; mutation; diagnosis; detection;
BMD; age-related macular dystrophy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human and mouse polynucleotides encoding CGICE polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human endothelin-1 antisense oligonucleotide fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 12.8; DB 1; 1
Pred. No. 2.9e+02;
n. Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 1 A; 6 C; 3 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Metzker M,
                                                                                                    Human CGICE PCR primer SEQ ID NO:8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 17; 67pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      269 GCCTGATCAAAGAGGA 284
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%;
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                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                            CO INC
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(UYUP-) UNIV UPPSALA.
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                                                                                                                                                                                                                          Homo sapiens
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18-DEC-1998;
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                                                               22-NOV-1999
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                                                                                                                                                                                                       Synthetic
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                          AAZ21232;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention provides antisense compounds of 8-30 nucleotides that inhibit the expression of human Cellular Inhibitor of Apoptosis-2 (c-1AP-2). The antisense compounds may be used for diagnostics, therapeutics (for modulating the expression of c-1AP-2), prophylaxis (e.g. to prevent or delay infection, inflammation, or tumor formation, as research reagents (e.g. to distinguish between members of a biological pathway) and in kits Sequences AAZ22103-142 represent phosphorothhoate ollopouncleotides used for antisense inhibition of cellular inhibitor of apoptosis-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense compounds complementary to Cellular Inhibitor of Apoptosis-2 useful for e.g. diagnostics, therapeutics, and as research reagents -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cellular Inhibitor of Apoptosis-2; antisense; diagnostic; therapeutic; c-IAP-2; prophylaxis; infection; inflammation; tumor formation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                            Gaps
AAZ41220, and AAY52701 to AAY52706, represent sequences used in the exemplification of the present invention.
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                                                                                                       Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Human c-IAP-2 mRNA inhibiting antisense oligo ISIS #23438
                                                                                                                                            Indels
                                                                                                    Score 12.8; DB 1;
Pred. No. 2.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 1 A; 6 C; 4 G; 7 T; 0 other;
                                                             Sequence 18 BP; 1 A; 6 C; 4 G; 7 T; 0 other;
                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cowsert LM, Ackermann EJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 3; Column 39; 33pp; English.
                                                                                                                                                                                                                                                                                                                         BP
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                                                                                                    Query Match 0.9%;
Best Local Similarity 87.5%;
Matches 14; Conservative
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AAZ21232/c
ID AAZ21232 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                       AAZ22129 standard; DNA; 18
                                                                                                                                                                                                           ATCTCCTGGGCTGTCT
                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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Gaps

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Length 18; Indels

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inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, acute asthma, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukemias, lymphonas, carcinomas e.g. colon cancer, breast cancer, lung cancer, pancreatic cancer, hepatocellular carcinoma, kidney cancer, melanoma, hepatic metastases, as well as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides may be derived from sequences AAX5272-74. These multiple target antisense treatment of diseases and conditions. Typical diseases and conditions are those associated with impaired respiration and
colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis; prostate cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense oligonucleotides used in treatment of, e.g. pulmonary vasoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 58; 120pp; English.
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                                                                                                                                                                                     Synthetic.
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    SYNCH COURT COURT
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0
                                                               Gaps
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                            0.9%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 2.9e+02; Live 0; Mismatches 2; Indels
Sequence 18 BP; 0 A; 4 C; 3 G; 11 T; 0 other;
                                                             Conservative
                                            Local Similarity
                                                           14;
                               Query Match
                                                Best Loca
Matches
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1143 CTTTTTTTTTTGG 1158 Cricirgicriring 16 à 음

AAX54193 standard; DNA; 18 BP. RESULT 618 AAX54193

AAX54193;

(first entry) 05-JUL-1999

Human endothelin-1 antisense oligonucleotide fragment.

Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; MAMAKAKEKEKEKEKE

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The specification describes antisense oligonucleotides (AAX52869-X55271)

directed against at least 2 mRNAs selected from target genes, coding and non-coding regions of RNAs corresponding to target genes, coding and concoding regions of the sequence of intiation codons, genomic flanking regions, intron-exon borders, the initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding protectins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides capecifically AAX5180-271 on be used for the capprocleotides (specifically AAX5180-271) can be used for the cancer conditions are those associated with impaired respiration and including lung diseases, pulmonary vasoconstriction, impeded inflammation, including lung diseases, pulmonary vasoconstriction, conditions, allergic rhinitis, acute ashima, allergies, asthma, impeded the sepiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary disease (COPD), and cancers such as leukemias, carcinomas e.g. colon cancer, breast cancer, lung cancer, pancrearic cancer, hepatocellular carcinoma, which may metastasize the particular sail as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer.
pulmonary hypertension; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; brast cancer; lung cancer; pancreatic cancer; hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Replication-competent; Sabin type 1 poliovirus vector; cloning site; 3C-protease cleavage site; mucosal vaccine; infectious disease; AIDS; human immunodeficiency virus type 1; HIV-1; small pox; poliomyelitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides used in treatment of, e.g. pulmonary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%; Score 12.8; DB 1; 1
ilarity 87.5%; Pred. No. 2.9e+02;
Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 0 A; 4 C; 3 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 57; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTTTTTTTTTTTGG 1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cricrigicitities 16
                                                                                                                                                                                                                                                                          98US-0093972.
                                                                                                                                                                                                                                       98WO-US19419
                                                                                                                                                                                                                                                                                                                                     (UYEC-) UNIV EAST CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX26160 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Primer for cDNA synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                 prostate cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-229400/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vasoconstriction
                                                                                                                                                                                                                                                                          09-JUN-1998;
17-SEP-1997;
                                                                                                                                                                                                                                       17-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-MAY-1999
                                                                                                                                                        WO9913886-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14;
                                                                                                                                                                                                 25-MAR-1999.
                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX26160;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1143
                                                                                                                                                                                                                                                                                                                                                                              Nyce JW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 619
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Cohen D, Blumenfeld M,
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   21-MAY-1997;
31-JUL-1997;
28-NOV-1997;
02-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9954500-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-OCT-1999
                                                                                                                                                                                                                                                                                                                    Vh and Vk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ73148;
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                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                          Carr FJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 621
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                                                                                                                                                                                                                                                         The invention relates to a replication-competent recombinant Sabin type 1 poliovirus vector encoding a multiple cloning site and 3C-protease cleavage site between the two end N-terminal residues. This comprises a vector containing an exogenous vaccine gene at the multiple cloning site. A method of production of both vectors is also provided. The recombinant number of infectious diseases, including human immunodeficiency virus type 1 (HIV-1) (which causes acquired immunodeficiency syndrome (AIDS)), small pox, poliomyelitis and Hepatitis C. The poliovirus-mediated mucosal vaccine vectors overcome the diseavantages exhibited by Mahoney vectors by being safe to humans, replicable (having equal replication ability to that of the wild type) vectors, where the introduced vaccine genes are
Hepatitis C; acquired immunodeficiency syndrome; Mahoney vector; viral;
                                                                                                                                                                                               New replication-competent recombinant Sabin type 1 poliovirus vector - useful for developing mucosal vaccines against \rm HIV-type 1, small pox, poliomyelitis and hepatitis C
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Non-immunogenic; epitope; T-cell; immunogenicity; immune system; SK; immunoglobulin; therapeutic; streptokinase; de-immunised; 708;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ·,
                                                                                                                                                                                                                                                                                                                                                                                                                               tch 0.9%; Score 12.8; DB 1; Length 18; al Similarity 87.5%; Pred. No. 2.9e+02; 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            708 Vk constructing flanking primer DIVX7.
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 1 A; 10 C; 4 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                       stably maintained during viral passages.
                                                                                                                                                                                                                                        Disclosure; Page 58; 64pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        302 CTGTGGGGGCTGCAAC 317
                                                                                           98WO-KR00242.
                                                                                                                97KR-0037812
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                                                                                                                                    (ALTW-) ALTWELL BIOTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV81061 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17 CGGTGGGGGGGGCAAC
                                                                                                                                                                            WPI; 1999-167434/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                          Jung HR;
                                                   WO9907859-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            De-immunised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9852976-A1
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                                                                                                                07-AUG-1997;
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                                                                       18-FEB-1999
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            primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer; ss
                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV81061;
                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                       Bae YS,
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proteins. The method comprises determining at least part of the amino acid sequence of the protein; (b) identifying in the amino acid sequence one or more potential epitopes for T-cell epitopes) of the given species; and (c) modifying the amino acid sequence to eliminate at least one of the T-cell epitopes identified in step (b) thereby to eliminate or reduce the immunogenicity of the protein when exposed to the immune system of the given species. A method of analysing a pre-existing protein to predict the basis for immunogenic responses is also provided. The methods can be used particularly for reducing the immunogenicity of immunoglobulins or therapeutic proteins, e.g. Streptokinase (SX). The products can be used for diagnosis and therapy. Sequences AAV81047-68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a method for the production of non-immunogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                             Reducing immunogenicity of proteins - by modifying the amino acid sequence of the protein to eliminate potential epitopes for T-cells of a given species
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human biallelic marker upstream amplification primer SEQ ID NO:7504.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human genome, biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyphing; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%; Score 12.8; DB 1;
87.5%; Pred. No. 2.9e+02;
ative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 2 A; 5 C; 4 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; Fig 16; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           991 TTCAGATCCGGCTTGG 1006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                      97GB-0016197.
97GB-0025270.
97US-0067235.
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97GB-0010480.
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98US-0109732.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                           (BIOV-) BIOVATION LTD
                                                                                                                                                                                                                                                                                                                        WPI; 1999-045301/04.
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AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ6579 to AAZ740 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmacceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaccutical agents acting on a disease as well as other
                                                                biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N.B. The SEQ ID NOs 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the present invention.
                                                                                                                                          Claim 9; Page 1830; 2745pp; English.
                                                                                            map of the human genome
                    WPI; 2000-013267/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           treatment.
                                                                  Novel
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Gaps .. 0 Query Match 0.9%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 2.9e+02; Matches 14; Conservative 0; Mismatches 2; Indels Sequence 18 BP; 4 A; 4 C; 4 G; 6 T; 0 other; 865 3 TCAGCATAACGCTTGG 18 850 TCAGCATACCGCTTTG à d

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AAZ76819 standard; DNA; 18 BP. (first entry) 10-SEP-2001 AAZ76819; RESULT 622 AAZ76819 

Human biallelic marker downstream amplification primer SEQ ID NO:11175.

Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; SNP; PCR primer; diagnosis; ss.

Homo sapiens.

WO9954500-A2

28-OCT-1999

99WO-IB00822. 21-APR-1999;

98US-0082614. 98US-0109732. 23-NOV-1998; 21-APR-1998;

(GEST ) GENSET

Chumakov I; Blumenfeld M, Cohen D,

WPI; 2000-013267/01.

Novel biallelic markers used to construct a high density disequilibrium map of the human genome

Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions -

WPI; 2000-679539/66.

Nyce JW;

Claim 14; Page 241; 1592pp; English

Claim 9; Page 2613; 2745pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection; chronic obstructive pulmonary disease; pulmonary infection; bronchitis;
           AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to PAZ7740 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of
                                                                                                                                                                   pharmaceutical agents and diagnostic methods, as well as the charmacterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other
                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                             N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the present invention.
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                                                                                                                                                                                                                                                                                                                                                        0.9%; Score 12.8; DB 1; Length 18;
17.5%; Pred. No. 2.9e+02;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human endothelin-1 polynucleotide fragment #1326.
                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 8 A; 5 C; 4 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                955 AGACTGCAGGACTGAC 970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF19759 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 ACACAGCAGGACTGAC 18
                                                                                                                                                                                                                                                                                                                                                                           87.58;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-MAR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 87.5'
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer; ss.
                                                                                                                                                                                                                                  treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF19759;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 623
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us09904568-3.rng

The present invention describes low adenosine (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides and compositions (I) compared to alternative base. (I) can have respiratory, bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. The antisense oligonucleotides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with lung/respiratory disorders and malignancies, such as stimulating and cativity of target polypeptides associated with a lung/respiratory disorders and malignancies, such as stimulating and chief receptors and antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, immunoglobulins and antibody receptors, bradykinin receptors, entral chemokines, adenosine receptors, bradykinin receptors, central nervous system (CKS) and peripheral nervous and non-nervous system peptide transmitters, defensine, growth factors, vasoactive peptides and receptors, bradykinin receptors, bradys proteins and malignancy associated proteins. The antisense oligonucleotides may be used in this way to treat disorders including respiratory obstruction (especially pulmonary obstruction and/or bronchoconstriction) and/or lung inflammation, allergic system and proteins as and malignancy associated with a disease or condition selected from pulmonary vasoconstriction, inflammation, allergic system (CKD), pain, cystic fibrosis (CF), allergic rhintis (RR), pulmonary hypertension, emphysema, chronic obstructive pulmonary infections, bronchitis, and/or cancer Abrila to Abrila are present hymen production. and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplification of the present invention. 

Sequence 18 BP; 0 A; 4 C; 3 G; 11 T; 0 other;

ö Gaps ; 0 0.9%; Score 12.8; DB 1; Length 18; 77.5%; Pred. No. 2.9e+02; ve 0; Mismatches 2; Indels 87.5%; Query Match 0.9 Best Local Similarity 87.5 Matches 14; Conservative

1143 CTTTTTTTTTTGG 1158 16 1 cricirgicirities à 용

RESULT 624 AAF19769

AAF19769 standard; DNA; 18 BP. AAF19769; 

(first entry) 14-MAR-2001

Human endothelin-1 polynucleotide fragment #1336.

Low adenosine antisense oligonucleotide, phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension, emphysema, pulmonary transplantation rejection, chronic obstructive pulmonary disease; pulmonary infection; bronchitis; cancer; ss.

Homo sapiens.

WO200062736-A2.

26-OCT-2000.

24-MAR-2000; 2000WO-US08020.

99US-0127958 06-APR-1999; (UYEC-) UNIV EAST CAROLINA.

(NYCE/) NYCE J

Nyce JW;

WPI; 2000-679539/66.

Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions -

Claim 14; Page 242; 1592pp; English.

in present invention describes for adenosities (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base. (I) can have respiratory, bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiathmatic, hypotensive and cytostatic activities. The antisense oligonucleotides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with ung/respiratory disorders and atribody polypeptides associated with activating peptide factors and transmitters, transcription factors, immunoglobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, binding proteins, adhesion molecules and their receptors, cytokine and chemokine receptors, adenosine receptors, bradykinin receptors, central nervous system (CNS) and peripheral nervous and non-nervous system receptors, binding proteins and malignancy associated proteins. The antisense oligonucleotides may be used in this way to treat disorders including respiratory obstruction (especially pulmonary obstruction and/or bronchoonstriction) and/or lung inflammation, allergy(ies) and/or bronchoonstriction) and/or bronchoonstriction) and/or lung inflammation, allergy(ies) and/or surfactant hypoproduction which are associated with a disease or condition selected from pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiratory pulmonary disease (COPD), pain, cystic fibrosis (CP), allergic rhinitis (RN), pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), pain, cystic fibrosis (CP), allergic rhinitis (RN), pulmonary and/or candor candor and/or and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplification of content antisense The present invention describes low adenosine (A)

Sequence 18 BP; 0 A; 4 C; 3 G; 11 T; 0 other;

Gaps .; 0 Length 18; Indels 0.9%; Score 12.8; DB 1; ilarity 87.5%; Pred. No. 2.9e+02; Conservative 0; Mismatches 2; Local Similarity ses 14; Conserv Query Match Matches

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Dp

à

RESULT 625

BP AAA86618/c ID AAA86618 standard; DNA; 18

AAA86618;

04-DEC-2000 (first entry)

Cdc 2 kinase hammerhead ribozyme recognitoin site #49.

Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.

Mammalia.

WO200032765-A2. 

08-JUN-2000.

99WO-US28772. 06-DEC-1999; 0

Gaps

0;

Indels

Score 12.8; DB 1; Pred. No. 2.9e+02; 0; Mismatches 2;

0, 615

0.9%;

BP.

Length 18;

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The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDKI, PCNA and Cyclin B1
                                                                                                                                                                                                                                                                                                                                                                                               Ribozyme; hairpin; hammerhead; gene therapy; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                             Cdc 2 kinase hammerhead ribozyme recognitoin site #50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 19; 109pp; English.
                                                                                                                                                                                                                                                              600 CAGCCTGAAGCCTGAC 615
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99WO-US28772
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98US-0110954
                                                                                                                                                                                                                                                                               cárccrgaagacrgac
                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Welch PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-412314/35.
                                                                                                                                                                                                                                                                                                                          AAA86619 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200032765-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-DEC-1999;
        04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                             04-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                        restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-JJN-2000
                                                                                                                                                                                                                                                                               18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rritz R,
                                                                                                                                                                                                                                                                                                                                            AAA86619
                                                                                                                                                                                                                                                                                                                                                                                                                         Mammalia
                                          Tritz R,
                                                                                                                                                                                                                                                                                                        RESULT 626
                                                                                                                                                                                                                                                                                                                  AAA86619/
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BP.

DNA; 18

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Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restrenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in restenosis treatment.
                                                                                                                                                                   Sequence 18 BP; 3 A; 3 C; 6 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA29110 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-SEP-2000 (first entry)
                                                                                                                                                                                                        Query Match
Best Local Similarity 87.59
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                               600 CAGCCTGAAGCCTGAC
                                                                                                                                                                                                                                                                                                                                                                                    CATCCTGAAGACTGAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA29110;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 62
AAA29110/
        888888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    o;
                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1.
Representative examples of ribozyme recognition sites are given in AAA8215 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme is resistant to endonuclease activity and hence is efficient in restenosis treatment.
                                                                                                                                                                                                                                             New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.9%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 2.9e+02; iive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 4 A; 3 C; 5 G; 6 T; 0 other;
                                                                                                                                       Robbins JM;
                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 19; 109pp; English.
                                                                                                                                       Barber JR,
                               98US-0110954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.9
Best Local Similarity 87.5
Matches 14; Conservative
                                                                               (IMMU-) IMMUSOL INC
                                                                                                                                                                                          WPI; 2000-412314/35
                                                                                                                                       Welch PJ,
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AAA29110-11 are primers, which can be used to clone an eosinophil-specific promoter (e.g. AAA29109) from the 5' flanking sequences of genomic cosinophil peroxidase gene. The promoter is useful for manipulating expression of heterologous proteins such as murine or human major basic proteins, eosinophil peroxidase, human granule ribonuclease, ribozyme, cell toxin and reporter proteins such as beta-galactosidase, within eosinophils. Transgenic animals are useful producing large quantities of eosinophils and as animal models for pulmonary pathologies such as asthma, using eosinophil-specific expression of transgenes. Discovery of the promoter is advantageous because eosinophil-specific expression of proteins is not toxic to the eosinophil and absence of eosinophils does not affect survival of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleotide comprising an eosinophil-specific promoter for expression of proteins such as cell toxins or eosinophil peroxidase in eosinophils and for ablating the eosinophil lineage from transgenic
                                                                                                                     Bosinophil; promoter; peroxidase; heterologous protein expression;
major basic protein; granule ribonuclease; transgenic animal model;
tissue-specific; primer; ss.
                                                                                        Primer 1 for eosinophil peroxidase gene promoter amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                     (MAYO-) MAYO FOUND MEDICAL EDUCATION & RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure, Page 6, 40pp, English.
                                                                                                                                                                                                                                                                                                                                                                                                                         Lee NA, Macias MP;
                                                                                                                                                                                                                                                                                                               99WO-US29162
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-42336B/36.
                                                                                                                                                                                                                                      WO200034304-A1
                                                                                                                                                                                                   Mus musculus.
                                                                                                                                                                                                                                                                                                             09-DEC-1999;
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Gaps

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Length 18; Indels

ch 0.9%; Score 12.8; DB 1; 1 1 Similarity 87.5%; Pred. No. 2.9e+02; 14; Conservative 0; Mismatches 2;

Query Match Best Local Similarity Matches 14; Conserv

Robbins JM;

Barber JR,

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CCAGCTCCAGGAGCTC 642
               ccadcrccaddaarc 2
627
ò
                8
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628

AAA12057 standard; DNA; 18 BP

AAA12057;

(first entry) 14-AUG-2000 Murine promoter OBhrel D4 HRE binding site primer.

HRE; hypoxia response element; hypoxia-inducible factor; HIF; vasotropic; caddiant; cytosfethic; antiathritic; gene therapy; ischaemia; arthritis; cardiovascular disease; peripheral arterial disease; cancer; primer phosphoglycerate kinase; PGK, muxine; promoter; OBhrel; ss.

Mus sp

WO200017371-A1.

30-MAR-2000,

99WO-GB03181 22-SEP-1999; 98WO-GB02885. 23-SEP-1998;

99GB-0001906. 99GB-0003538. 16-FEB-1999; (OXFO-) OXFORD BIOMEDICA UK LID.

Binley KM, Naylor S;

WPI; 2000-283595/24.

Novel polynucleotide constructs comprising at least two repeats of a hypoxia response element useful for driving expression of nucleic acids of interest in a cell under hypoxic conditions

Example 21; Page 109; 155pp; English.

This invention describes novel polymuclectide comprising at least 2 repeats of a hypoxia response element (HRE), where the hypoxia-inducible factor (HIE) consensus binding sites within each of the 2 repeats are separated by a spacer of at least 20 contiguous nucleotides. The products of the invention have vasotropic, cardiant, cytostatic and antiarthritic activity and can be used for gene therapy. The polymucleotides are useful for delivering nucleic acids of interest to mammalian cells. Lentiviral vectors are responsive to hypoxic agents and to agents that mind: vectors are responsive to hypoxic agents and to eagents the mind: typoxia. This regulation can be harnessed in vitro to enhance the production of the vector and can be used in vitro to regulate gene expression in response to a physiological signal. The vectors have utility in disease, where ischaemia, including hypoxia, is a feature, or gradiovascular disease, peripheral arterial disease, cancer and arthritis. The novel regulatory construct is capable of driving very high levels of transcription under normal oxygen conditions. The polynucleotide construct targets cells within a tumor mass that are under conditions of hypoxia without affecting normal surrounding tissue. This sequence represents a murine phosphoglycerate kinase (PGK) promoter OBhrel HRE binding site primer which is described in the method of the 

Sequence 18 BP; 3 A; 4 C; 8 G; 3 T; 0 other;

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Gaps
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 Length 18;
Best Local Similarity
Matches 14; Conserv
Query Match
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1235 TGGTGCTGGACGTGGC 1250
                             2 resrecaseacerese 17
                          DP
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RESULT 629

AAA33637 standard; DNA; 18 

AAA33637;

28-JUL-2000

(first entry)

Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothioate; impaired respiration; inflammation; allergy; allergy; disease; bronchoconstriction; inhibitor; antiinflammatory; antiallergic, attiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD; cancer; leukaemia; lymphoma; carcinoma; metastasis; ss. Low adenosine antisense oligonucleotide SEQ ID NO:1326.

Homo sapiens.

WO200009525-A2.

24-FEB-2000

99WO-US17712. 03-AUG-1999; 98US-0095212. 03-AUG-1998;

(UYEC-) UNIV EAST CAROLINA.

Nyce JW;

WPI; 2000-205971/18.

New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or cancers

Claim 18; Page 430; 1343pp; English.

The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antiallergies, and/or inflammation. The ON can have antiinflammatory, antiallergies, useful for the treatment of diseases associated with inflammation.

"Useful for the treatment of diseases and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary bypertension, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, carcinomas, and cancers such as leukaemias, lymphomas, carcinomas, and cancers such as leukaemias, lymphomas, breast and prostate cancer. The reduction of the adenosine content of the ONs reduces side effects. The A-containing ONS break down with the relacence of expediences which activates adenosine receptors causing bronchoconstriction and inflammation. AAA22313 to AAA33212 represent the nucleotide sequences given in the sequence listing from the present invention, which correspond to SED ID No:1 to 188, but the sequences differ from the previously named sequences. SEQ ID No:11 to 160 (AAA32321 to AAA32322) to AAA33232 to AAA33232 to AAA33232 to and then the last invention do not match we with their corresponding SEQ ID No: sequences given in the sequence listing. Corresponding SEQ ID No: sequences given in the sequence listing.

Page 310

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The present invention describes a new composition comprising an antisense oligomucleotide (OM) with low adenosine (up to 15%), which targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antiallergies, and/or inflammation. The ON can have antiinflammatory, antiallergies, and/or inflammation. The Compositions are useful for the treatment of diseases associated with inflammation, impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, carcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the ONs reduces side effects. The A-containing ONs break down with the release of decoxyadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA35313 to AAA55312 represent invention, which correspond to SEQ ID No:1 to 185, but the sequences
                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                       Human, adenosine receptor; low adenosine antisense oligonucleotide; phosphorothicate; impaired respiration; inflammaton; allergy; allergic disease; bronchoconstriction; inhibitor; antinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chonic obstructive pulmonary disease; COPD;
                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                ;
                                        Length 18;
                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.
                                                                                                                                                                                                                                                                                                                                                                                   Low adenosine antisense oligonucleotide SEQ ID NO:1336.
                                   Score 12.8; DB 1;
Pred. No. 2.9e+02;
0; Mismatches 2;
Sequence 18 BP; 0 A; 4 C; 3 G; 11 T; 0 other;
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                                                                                                                        1143 CTTTTTTTTTGG 1158
                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                            cricrigiciriride 16
                                     ch
1. Similarity 87.5%;
14; Conservative
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                                                                                                                                                                                                                                                               AAA33647 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                          Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                     AAA33647;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tissue relating to obesity, particularly complications of visceral obesity including diabetes, hyperlipemia, hypertension, arteriosclerosis, hyperuricemia and sleep apnea syndrome. The genes (AAZ96611-633) and the proteins (AAX67698-Y67600) are used in the genetic diagnosis, prevention and treatment of adipose tissue related diseases. Sequences AAZ90640-51 represent PCR primers amplifying the human adipose tissue genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to identification of genes and proteins of adipose
differ from the previously named sequences. SEQ ID NO:11 to 1680 (AAA33323 to AAA33992) are specifically claimed ONs from the present invention. N.B. Sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequences
                                                                                                                                                                                                                                                                                                                                                                                                           Adipose tissue, obesity, diabetes, hyperlipemia, hypertension, human, arteriosclerosis, hyperuricemia, sleep apnea syndrome; PCR primer; ss
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                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     physiologically active protein specifically derived from mammal
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                                                                                                                                                   Indels
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37.5%; Pred. No. 2.9e+02;
tve 0; Mismatches 2;
                                                                                                                   0.9%; Score 12.8; DB 1;
7.5%; Pred. No. 2.9e+02;
ve 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                 gene amplifying primer #8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 1 A; 0 C; 2 G; 15 T; 0 other;
                                                                                        Seguence 18 BP; 0 A; 4 C; 3 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Page 18; 50pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1144 TITITITICITITIGGA 1159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                 1143 CTTTTTTTTTTGG 1158
                                                             given in the sequence listing.
                                                                                                                                                                                                                                                                                          AAZ90647 standard; DNA; 18 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98JP-0225228.
                                                                                                                                      87.5%;
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                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NISB ) JAPAN TOBACCO INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                     14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                 Human adipose tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-306578/27.
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Les 14; Conserv
                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-JUL-1998;
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                                                                                                                                                                                                                                                                                                                        AAZ90647;
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                                                                                                                       Query Match
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                                                                                                                                                      Matches
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The invention relates to antisense compounds, especially oligomucleotides, which are targeted to a nucleic acid encoding inducible oligomucleotides, which are targeted to a nucleic acid encoding inducible contervations of inducible nitric oxide synthase. The antisense compounds carportective, disorder and vascitopic activity. The antisense compounds control cective, disorder and vascitopic activity. The antisense coligomucleotides are useful for inhibiting the expression of inducible nitric oxide synthase in cells or tissues. In particular, the antisense coligomucleotides are useful for treating diseases or disorders associated with inducible nitric oxide synthase, e.g. diabetes, immunological disorder, cardiovascular disorder, neurological disorder or schaemia/reperfusion injury. The antisense oligomucleotides are also useful for research and disonseits. The present sequence is that of an antisense 2.O-methoxyethyl agamer oligomucleotide with a customorphicate backbone, a central "gap" region of ten nucleotides cresidues in the 2.-MOE wings are 5.-methoxyethyl) wings (cytidine cresidues in the 2.-MOE wings are 5.-methylcytidines) and targeted to human creating control of the control of the control of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 4 A; 8 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; Page 83; 144pp; English.
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99US-0161532.

26-OCT-1999;

Robbins JM, Tritz R; (IMMU-) IMMUSOL INC.

WPI; 2001-300427/31.

26-OCT-2000; 2000WO-US29500.

WO200130362-A2.

03-MAY-2001.

Homo sapiens.

Synthetic.

/note = "phosphorothioate backbone, 5' and 3' four nucleotide 2'-MOE (2'-0-methoxyethyl) wings (the cytidine residues in the 2'-MOE wings are 5-methylcytidines) and a deoxy gap" New antisense oligonucleotides for modulating the expression of inducible nitric oxide synthase in cells or tissues, particularly useful for treating e.g. immunological, cardiovascular or neurological disorders, or ischaemia Antisense oligonucleotide; inducible nitric oxide synthase; NOS; modulate expression; immunomodulator; antidiabetic; cardiovascular; cardiant; neuroprotective; vasotropic; ischaemia; reperfusion injury; 2'-0-methoxyethyl; phosphorothioate; human; ss. Human inducible NOS antisense oligonucleotide SEQ ID NO 38. Location/Qualifiers ...18 /\*tag= a /mod base= OTHER Cowsert 15-JAN-2001; 2001WO-US01381 24-JAN-2000; 2000US-0490208 (first entry) (ISIS-) ISIS PHARM INC. Dean NM, WPI; 2001-465340/50. WO200152902-A1. modified base Homo sapiens 02-OCT-2001 26-JUL-2001. Bennett CF, AAH47994;

Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site, eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolyric; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.

Cdc 2 kinase hammerhead ribozyme recognition site SEQ ID NO:4208.

(first entry)

10-SEP-2001

AAH61784;

AAH61784 standard; DNA; 18

The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (1) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (1). (1) can have antipsoriatic, ophthalmological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (1) can be used in gene therapy. (1) and (II) are useful for treating proliferative skin diseases such as psoriaeis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, and for treating and preventing such as kelolid, adhesion and hypertrophic or hypertrophic burn and contract of the state of t Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent scarring such as keloid, adhesion and hypertrophic or hyper scar. AAH57577 to AAH62099 represent sequences used in the exemplification of the present invention. Disclosure; Page 379; 408pp; English. kinases 

Sequence 18 BP; 4 A; 3 C; 5 G; 6 T; 0 other;

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Gaps

.; 0

0.9%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 2.9e+02; tive 0; Mismatches 2; Indels

1239 GCTGGACGTGGCCATG 1254

à g

Local Similarity 87.5 nes 14; Conservative

Matches

Query Match

GCTTGAGGTGGCCATG 3

18

Gaps . 0 Length 18; Indels 2; 0.9%; SCOIE 12.8; DB 1; 87.5%; Pred. No. 2.9e+02; tive 0; Mismatches 2; Query Match Best Local Similarity 87.5°

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600 CAGCCTGAAGCCTGAC 615

6

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Human, ribozyme therapy, hairpin ribozyme, hammerhead ribozyme, recognition site, target, ribozyme binding site; eye disease, vulnerary; proliferative disease, skin disease, sociaais, diabetic retinopathy; cytokine; inflammation, cell-cycle dependent kinase, cyclin, MWP; matrix metalloproceinase, growth factor, reductase, scarring; cytostatic, antipsoriatic; dermatological; antisebornheic; antidiabetic; virucide, antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis, actinic keratosis, squamous cell carcinoma; basal cell carcinoma; sebornheic wart; vitreoretinopathy; scar;
                                                                                                                Cdc 2 kinase hammerhead ribozyme recognition site SEQ ID NO:4209.
                                                    AAH61785 standard; DNA; 18 BP
                                                                                                                                                                                                                                 sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                   99US-0161532.
                                                                                                                                                                                                                                                                                                                               26-OCT-2000; 2000WO-US29500.
                                                                                            (first entry)
CATCCTGAAGACTGAC
                                                                                                                                                                                                                                                                                                                                                                                            Tritz R;
                                                                                                                                                                                                                                                                                                                                                                       (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-300427/31.
                                                                                                                                                                                                                                                                                     WO200130362-A2.
                                                                                                                                                                                                                                                     sapiens.
                                                                                                                                                                                                                                                                                                                                                   26-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                            Robbins JM,
                                                                                            10-SEP-2001
                                                                                                                                                                                                                                                                                                          03-MAY-2001
                                                                                                                                                                                                                                                                  Synthetia.
18
                                                                       AAH61785;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    kinases
                              RESULT 634
AAH61785/C
                                                                                                                                                                                                                                                        Homo
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New cloned Corynebacterium glutamicum thrE gene useful for producing thrE-overexpressing coryneform bacteria for production of L-threonine

Example 2; Page 7; 22pp; German.

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Sahm

Eggeling L,

Ziegler P,

Thierbach G,

WPI; 2001-227606/24.

(DEGS ) DEGUSSA-HUELS AG. (KERJ ) FORSCHUNGSZENTRUM JUELICH GMBH.

99DE-1041478. 99DE-1041478.

01-SEP-1999; 01-SEP-1999;

08-MAR-2001.

C. glutamicum ATCC13032 thrE PCR primer thrE-reverse. L-threonine; thrE; animal nutrition; human medicine;

(first entry)

25-MAY-2001

AAF61243;

AAF61243 standard; DNA; 18 BP

RESULT 635

AAF61243,

615

600 CAGCCTGAAGCCTGAC

16 CATCCTGAAGACTGAC 1

엄 à

pharmaceutical industry; PCR primer; ss.

Corynebacterium glutamicum.

DE19941478-A1.

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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (1) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (WMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, ophthalmological, cytostatic, antiseborrheic, antidiabetic, antisickling, captubalmological, vulnerary, keraclotytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriatis, atopic dermatitis, atopic dermaticis, atopic dermaticis, atopic dermaticis.

Skin diseases such as psoriatis, atopic dermatitis, atopic dermaticis.

Skin diseases such as psoriatis, atopic dermatitis, atopic dermaticis, atopic dermaticis, atopic dermatosis, atoming the cannopathy, vireoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adheeion and hypertrophic or hypertrophic burn sequences used in the
Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 379; 408pp; English.
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0;
                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel Corynebacterium glutamicum thrE gene (I). The invention also describes (1) an amino acid sequence that is derived from the nucleic acid sequence of (I) and is selected from two sequences of 489 amino acid sidven in the specification; (2) coryneform microorganisms transformed with (I); (3) production of L-threonine by culturing coryneform bacteria in which nucleotide sequences encoding the thrE gene (sic) are overexpressed; and (A) a process for producing (I). Coryneform bacteria that overexpress (I) are useful for producing L-threonine, which is useful in animal nutrition, human medicine and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.9%; Score 12.8; DB 1; Length 1
87.5%; Pred. No. 2.9e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human Akt-3 antisense oligonucleotide, SEQ ID NO: 85.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence 18 BP; 0 A; 6 C; 5 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    278 AAGAGGAAGCAGCAGC 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAGAGGAAACCGCAGC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF79677 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pharmaceutical industry.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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Gaps

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Length 18; Indels

Score 12.8; DB 1; Pred. No. 2.9e+02; 0; Mismatches 2;

0.9%;

Query Match 0.9 Best Local Similarity 87.5 Matches 14; Conservative

Sequence 18 BP; 3 A; 3 C; 6 G; 6 T; 0 other;

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WPI; 2001-147509/15.
                                                                                                              WPI; 2001-147509/15.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shimkets RA,
11-AUG-1999;
10-AUG-2000;
                                                                              Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-FEB-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF74483
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                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is one of a number of antisense compounds of up to 30 nucleobases in length targeted to a nucleic acid encoding human Akt-3. The antisense compounds are useful for inhibiting the expression of human Akt-3 in human cells or tissues. They are also useful for modulating the expression of Akt-3, and for treating a human or an animal suspected of having, or being prone to, a disease or condition associated with Akt-3 expression. The antisense compounds may also be used as research reagents, in kits and in diagnostics, e.g. to elucidate the function of a particular gene or to distinguish between functions of various members of
            Human, Akt-3; protein kinase; cytostatic; antiinflammatory; infection; antisense therapy; inflammation; tumour; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   biological pathway; and as a prophylactic, e.g. to prevent or delay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                             New antisense compounds targeting nucleic acids encoding human Akt-3 useful for treating a disease or condition associated with Akt-3 expression, or in preventing or delaying inflammation or tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.9%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 2.9e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Clone 21399247.0.1 PRO5 sequencing primer SEQ ID NO:66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 5 A; 5 C; 4 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    infection, inflammation or tumour formation.
                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Column 40; 37pp; English.
                                                                                                                                                                                                                                                              Roth RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF74480 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   375 CCAGCTTCCTCCAGAG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 ccadriracrccadag 17
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                                                                                                                                                                                              99US-0474922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                              Cowsert LM,
                                                                                                                                                                                                                                                                                               WPI; 2001-264979/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200110902-A2
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                                                                Homo sapiens
                                                                                             US6187586-B1
                                                                                                                                                             29-DEC-1999;
                                                                                                                                                                                              29-DEC-1999;
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                                                                                                                              13-FEB-2001,
                                                                                                                                                                                                                                                                                                                                                                                 formation
                                                                                                                                                                                                                                                              Monia BP,
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637 RESULT 63

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The present invention describes isolated nucleic acids encoding secreted polypeptides, designated PROX polypeptides (i.e. a PRO polypeptide where X is an integer from 1 to 17). PROX polypeptides have cytostatic, amunomodulatory and reproduction activities, and can be used in gene therapy, and as PROX antagonists and PROX agonists. PROX polypeptides, nucleic acids and antibodises are useful in the manufacture of a medicament for treating a syndrome associated with a PROX associated disorder, e.g. a cell proliferation and/or differentiation disorder (e.g. cancer or immune associated disorders) and a gestational disease (e.g. pre-clampsia). They are also used for screening for a modulator of activity or of latency or predisposition to a PROX-associated disorder. AMF74412 to AAF7448 encode the specifically claimed human PROX polypeptides PRO1 to PRO17 given in AAB70531 to AAB70547. The present section of an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human, PRO; PROX; cytostatic; immunomodulatory; reproduction;
gene therapy, cell proliferation; differentiation disorder; cancer;
immune associated disorder; gestational disease; pre-clampsia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                               Nucleic acids encoding secreted polypeptides, designated PROX polypeptides, useful for treating a syndrome associated with a PROX-associated disorder, e.g. cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.9%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 2.9e+02; ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clone 21399247.0.1 PROS sequencing primer SEQ ID NO:69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 4 A; 5 C; 7 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 9; Page 125; 166pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer; sequencing primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1292 TIGCICAGCCIGGCC 1307
99US-0148433.
2000US-0148433.
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10-AUG-2000; 2000US-0148433
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Matches 14; Conservative
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                                                                                                                   (CURA-) CURAGEN CORP.
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Sequence 18 BP; 2 A; 7 C; 5 G; 4 T; 0 other;
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The present invention describes isolated nucleic acids encoding secreted polypeptides, designated PROX polypeptides (i.e. a PRO polypeptide where is an integer from 1 to 17). PROX polypeptides have cytostatic, immunomodulatory and reproduction activities, and can be used in gene therapy, and as PROX antagonists and PROX agonists. PROX polypeptides, nucleic acids and antibodies are useful in the manufacture of a medicament for treating a syndrome associated with a PROX-associated disorder, e.g. a cell proliferation and/or differentiation disorder (e.g. pre-clampsia). They are also used for screening for a modulator of activity or of latency or predisposition to a PROX-associated disorder. PAF74432 to AAF74448 encode the specifically claimed human PROX polypeptides PRO1 to PRO17 given in AAB70531 to AAB70547. The present sequence represents a primer used in an example from the present
Nucleic acids encoding secreted polypeptides, designated PROX polypeptides, useful for treating a syndrome associated with a PROX-associated disorder, e.g. cancer -
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Gaps
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0
0.9%; Score 12.8; DB 1; Length 18; ilarity 87.5%; Pred. No. 2.9e+02; Conservative 0; Mismatches 2; Indels
                                                                     1292 TIGCICAGCCIGGCC 1307
                   Local Similarity
hes 14; Conserv
      Query Match
                    Best Loca
Matches
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AAF68906 standard; DNA; 18 (first entry) COXII probe #12 12-APR-2001 AAF68906; RESULT 639 AAF68906 

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Mitochondria; cytochrome C oxidase; COX; Alzheimer's disease; probe; ss.

Homo sapiens

US6171859-B1

09-JAN-2001

95US-0413740 30-MAR-1995;

94US-0219842 30-MAR-1994;

(MITO-) MITOKOR

Parker WD; Herrnstadt C,

WPI; 2001-136875/14.

cytochrome C oxidase activity for diagnosing Alzheimer's disease, involves contacting mitochondria with a conjugate of targeting molecule Targeting conjugate molecule to mitochondria having defective and

Disclosure; Columns 21-22; 88pp; English.

The present invention relates to a method for selectively accumulating a mitochondrial disabling or destructive amount of a conjugate molecule in mitochondria having defective cytochrome C oxidase (COX) activity or

Length 18;

Score 12.8; DB 1; Pred. No. 2.9e+02;

0.98; 87.5%;

Best Local Similarity

Query Match

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This invention describes novel isolated, preferably recombinant, polypeptides (I) containing the amino acid sequence for human procalciorain (hpcT). The producte of the invention have antitumor, antisepsis and antimilammatory activity. (I) (also antibodies (Ab) raised against it) are used: (i) for diagnosis and treatment of tumors, sepsis and systemic inflammatory response syndrome, (ii) to raise Ab; (iii) for quantitative or qualitative detection and analysis, especially of hpcT and antibodies against it; (iv) as controls or standards for assays; and (v) for affinity chromatography. Isolated (I) can be produced inexpensively in large amounts by recombinant expression. Solutions of (I) that contain a polyethoxylated sterol ester have good storage stability. This sequence represents a PCR primer used in the
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displaying increased membrane potential. The method involves contacting mitochondria with a conjugate molecule comprising a targeting molecule conjugated to a toxin, where the conjugate or targeting molecule selected accumulates in the mitochondria. The method is useful for diagnosis of Alzheimer's disease (AD), especially sporadic AD. The present sequence is a probe used in the method of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New, preferably recombinant, human procalcitonin, useful for diagnosis and treatment of sepsis, tumors and systemic inflammatory response
                                                                                                                                                                                                           Gaps
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                                                                                                                                                                   0.9%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 2.9e+02; ive 0; Mismatches 2; Indels
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                                                                                                                                  Sequence 18 BP; 3 A; 9 C; 3 G; 3 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human procalcitonin pCT PCR primer 1101
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03-APR-2000; 2000DE-1016278.
08-JUN-2000; 2000DE-1027954.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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The invention relates to isolated genes (Gene 216) from human chromosome 20p13-p12 and the proteins they encode. The nucleic acids and proteins may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate Gene 216 expression. For example, the nucleic acids (or vectors) and proteins may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of gene 216 by expressing in a patient's genome that affect the activity of gene 216 by expressing contactive proteins or to supplement the patients own production of Gene 216 proteins, by inserting the nucleic acids into a host cell and culturing the cell to express the protein. The nucleic acids assays to detect and quantitate the presence of similar nucleic acids assays to detect and quantitate the presence of similar nucleic acid sequences in samples and therefore which patients may be in need of ceptorative therapy. The Gene 216 protein may also be used as antigens in the production of antibodies against Gene 216 and in assays to identify modulators of Gene 216 expression and activity. The anti-Gene 216 antibodies may also be used as antigens of antibodies and the presence of Gene 216 and in assays to identify and additions of Gene 216 antibodies may also be used as diagnostic agents for detecting the presence of Gene 216 proteins in samples (e.g. procession and activity. The anti-Gene 216 antibodies may also be used as diagnostic agents for detecting the presence of Gene 216 proteins in samples (e.g. procession and activity. The anti-Gene 216 proteins in samples (e.g. procession and activity. The anti-Gene 216 proteins in samples (e.g. procession and activity. The anti-Gene 216 proteins in samples (e.g. procession and activity. The anti-Gene 216 proteins in samples (e.g. procession and activity. The protein and activity and any protein may also be used as diagnostic or agents for detecting the presence of Gene 216 proteins in samples (e.g. pro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, Gene 216, chromosome 20p13-p12, antiasthmatic; anorectic; antiinflammatory; gastrointestinal; gene therapy; vaccine; asthma; obesity; inflammatory bowel disease; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Isolated genes (Gene 216) from human chromosome 20p13-p12 and t
proteins they encode, useful for the prevention, diagnosis and
treatment of asthma, obesity and inflammatory bowel disease
2;
                                                                                                                                                                                                                                                                                                                                                                                                  Gene 216 SSCP sequencing primer SEQ ID NO 178.
Mismatches
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                                                                                                                                                                                                                                     ABZ72206 standard; DNA; 18
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                                                     833 TGAAGCTTTCAGATGG
                                                                                                       rgaagcrirragrigg
14; Conservative
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  Matches
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Sequence 18 BP; 2 A; 7 C; 3 G; 6 T; 0 other;

0; ; 0 The invention relates to a truncated neublastin polypeptide comprising an amino acid terminus that lacks one or more amino-terminal amino acids of a mature neublastin polypeptide. The polypeptides and nucleic acids are useful for treating neurodegenerative disorders such as ischemic neuronal damage, traumatic brain injury, peripheral neuropathy, neuropathic pain, Alzheimer's disease, Huntington's disease, Parkinson's disease, amyotrophic lateral sclerosis, memory impairment, diabetes, renal diseases, or glaucoma by moderating metabolism, growth, differentiation or survival of a nerve or neuronal cell. This polynucleotide sequence is Nootropic; neuroprotective; antiparkinsonian; anticonvulsant; analgesic; tranquiliser; antidiabetic; ophthalmological; neurodegenerative disorder; neublastin; ischemic neuronal damage; traumatic brain injury; diabetes; peripheral neuropathy; neuropathic pain; Alzheimer's disease; glaucoma; Huntington's disease; parkinson's disease; glaucoma; memory impairment; renal disease; PCR; primer; ss. New truncated neublastin polypeptides lacking one or more amino-terminal amino acids of a mature neublastin polypeptide useful for treating neurodegenerative disorders, e.g. peripheral neuropathy, neuropathic pain, brain injury -Gaps Gaps . 0 0; Length 18; Length 18; Indels Indels 0.9%; Score 12.8; DB 1; 37.5%; Pred. No. 2.9e+02; 1ve 0; Mismatches 2; Score 12.8; DB 1; Pred. No. 2.9e+02; ); Mismatches 2; Sequence 18 BP; 1 A; 6 C; 9 G; 2 T; 0 other; primer of the invention. Rossomando A; Neublastin DNA related PCR primer. Disclosure, Fig 8; 138pp, English. 0; 731 0.9%; 3 TGGCCCGGCTGCAGGG 18 87.5%; 12-MAR-2002; 2002WO-EP02691 12-MAR-2001; 2001US-0804615 ABT11916 standard; DNA; 18 19-DEC-2002 (first entry) r survival of a nerve or neublastin PCR primer of 716 TGGCCCAGCAGCAGGG 16 AGCAGAGGGCATGGCT Best Local Similarity 87.5 Matches 14; Conservative 14; Conservative Sah DWY, Johansen TE, WPI; 2002-713515/77. (BIOJ ) BIOGEN INC (NSGE-) NS GENE AS Query Match Best Local Similarity WO200272826-A2 Unidentified. 19-SEP-2002. 725 ABT11916; Query Match Matches RESULT ð 쉽 à d

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Reelin protein CR-50 epitope region, useful for diagnosis and treatment
                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an isolated acyl coenzyme A thioesterase (ACH) encoding nucleic acid, encoding one of ACH1, ACH2, ACH4 or ACH5. ACH enzymes have a role in lipid oxidation, regulation of Coenzyme A pools and in fatty acid metabolism. Also include are a host cell transfected with the nucleic acid, a transgenic plant transfected with the nucleic acid, a transgenic plant transfected with the nucleic acid is useful for altering a phenotype of a plant and for making a transgenic plant, by transfecting the plant tissue with the ACH nucleic acid under conditions such that a transgenic plant is generated. The ACH nucleic acid under conditions such that a transgenic plant is generated. The ACH nucleic acid is also useful for producing variants of acyl-CoA thioesterases. The present sequence is a PCR primer used to amplify Arabidopsis ACH4 encoding sequences.
                                                                                                                                                                                                                                                                                         Novel acyl coenzyme A thioesterase gene useful for altering a phenotype of a plant, making a transgenic plant and for producing variants of acyl-CoA thioesterases
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cerebral disturbance, reelin protein, neuroprotective, PCR primer; ss.
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                                                                                                                                                                                                                      Browse JA;
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14; Conservative
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ID ABL40174 standard; DNA; 18
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                                       19-JUL-2001; 2001WO-US22907.
                                                                                              16-JUL-2001; 2001US-0906408.
                                                                           21-JUL-2000; 2000US-220028P
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                                                                                                                                      (TILT/) TILTON G B. (SHOC/) SHOCKEY J M. (BROW/) BROWSE J A.
                                                                                                                                                                                                                                                          WPI; 2002-241573/29
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Best Local Similarity
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31-JAN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              involving taking a DNA-containing sample from an organism or cell culture that has been treated with a test compound and determining any changes in the DNA methylation status or pattern caused by said test compound. The method is used for diagnosis and prognosis of adverse toxic responses in individuals. The present sequence is a human sequence used to demonstrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Toxicological diagnosis, useful for diagnosis and prognosis of adverse reactions, based on effect of test compounds on methylation status of selected genes, involves determining changes in DNA methylation status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present invention relates to a method of toxicological diagnosis,
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                                                                                                                                                                                                     Toxicological diagnosis; DNA methylation; methylation status; toxic response; human; ds.
                                                                                                                                                               Human gene methylation status determination oligo SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.9%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 2.9e+02; cive 0; Mismatches 2; Indels
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                                                ABQ65394 standard; DNA; 18
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Matches 14; Conserv
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            RESULT 643
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glutamicum thrE gene.

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Example 2; Page 7; 16pp; Japanese.
                                                              Best Local Similarity
Matches 14; Conser
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The present invention describes the mouse reelin protein CR-50 epitope region, which contains the CR-50 antibody recognition site and is free from F-spondin domains and repetitive sites. Also described are: (1) an expression vector comprising a polymucleotide encoding a reelin protein epitope region; (2) host cells with transfected the expression vector; (3) polymeptides prepared by culture of the host cells; and (4) polymucleotides comprising the 351 base sequence given in AB1640165 which encodes the 117 amino acid sequence given in AB306244; and (5) use of the polymucleotide for diagnosis and/or treatment of diseases caused by abnormal positioning of neural cells, and stimulation of association of reelin protein. The mouse reelin protein CR-50 epitope region has neuroprotective activity, and can be used in the diagnosis and treatment of neurons. The present sequence represents a PCR primer for the mouse reelin protein CR-50 epitope region, which is used in an example from
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to the fermentative production of L-threonine using an Enterobacterium, particularly one already capable of producing L-threonine, in which activity of the thrE gene sequence (or sequences) is increased by overexpression. L-threonine is useful in animal nutrition, human medicine and the pharmaceutical industry. The present sequence is a PCR primer used to isolate the Corynebacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fermentative production of L-threonine, useful in animal nutrition, comprises culturing enterobacterium with increased thrE gene activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (AB10512-AB151809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals.
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87.5%; Pred. No. 2.9e+02;
ative 0; Mismatches 2; Indels
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Seguence 18 BP; 0 A; 6 C; 5 G; 7 T; 0 other;
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Homo sapiens
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Pred. No. 2.9e+02;
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Matches 14; Conservative
                                                                                                      (first entry)
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                           ABL30631/
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Human; human leukocyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.

Human HLA genotyping oligonucleotide SEQ ID NO 158.

(first entry)

21-MAR-2002

EXXXEXEXXXXXX

ABL30669;

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The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (ABL50512-ABL51809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of pancreas tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, langue for transplantation e.g. of bone marrow, kidney, liver, diagnosis of genetic diseases and identifying individuals.
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87.5%; Pred. No. 2.9e+02;
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                                                                                                            Ichihara T,
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01-JUN-2000; 2000JP-0164798.
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Local Similarity 87.5
Local 14, Conservative
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Nishida M;

Matsumura Y,

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Moriya S,
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Kagiya T,
                                                                                   WPI; 2002-122074/16
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.. 0 Indels Mismatches 0; 143 CGCTCGGCTCCGCTCC 158 cecrcedercererce Conservative Query Match Best Local Similarity 14; 18 Matches g

ABL31108 standard; DNA; 18 BP ABL31108; RESULT 651

Human HLA genotyping oligonucleotide SEQ ID NO 597. (first entry) 21-MAR-2002 

Human; human leukocyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.

Homo sapiens

WO200192572-A1

06-DEC-2001

01-JUN-2001; 2001WO-JP04662

01-JUN-2000; 2000JP-0164798

(NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.

Moriya S, Matsumura Y, Fi Ichihara WPI; 2002-122074/16. Kagiya T, Inoko H,

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Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when of individuals e.g. by detern transplanting between them -

Claim 10; Page 206; 345pp; Japanese.

The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (ABL30512-ABL31809) originating in the sequences of

0.9%; Score 12.8; DB 1; Length 18;

Sequence 18 BP; 3 A; 2 C; 7 G; 6 T; 0 other;

Query Match

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genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals.
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immunogenetic; transplantation; genetic disease; ss.
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87.5%; Pred. No. 2.9e+02;
tive 0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                      18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUN-2001; 2001WO-JP04662.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JUN-2000; 2000JP-0164798
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABL31109 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                    3 rcracedereacterr
                                                                                                                                                                                                                                                                           14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Inoko H, Kagiya T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-122074/16.
                                                                                                                                                                                                                                                         Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200192572-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL31109;
                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                     ABL31109
                                                                                                                                                                                                                                                                                                                                                                                                               RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                     8$333333333
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0
                                                                                                                                                          diabetes; obesity; infection; acromegaly; gigantism; sodium retention; water retention; metabolic syndrome; mood disorder; sleep disorder; Growth hormone dysfunction; familial growth hormone deficiency; short stature; pituitary storage defect; human; PCR primer; LCR25;
                                                                                                                                  Growth hormone 1 gene (GH1), locus control region (LCR) primer LCR25.
         Gaps
                                                                                                                                                   hormone 1; GH1; osteopathic; gene therapy; protein therapy;
         ô
Pred. No. 2.9e+02;
0; Mismatches 2; Indels
                          1182 TCTATAGGTGAGTGTT 1197
                                                                                    BP.
                                                                                                                                                                                              locus control region; LCR; ss
                                                                                                                                                                                                                                                                                         14-JUL-2000; 2000EP-0306004
                                                                                                                                                                                                                                                                14-MAY-2001; 2001WO-GB02126
                                                                                                                                                                                                                                                                                 12-MAY-2000; 2000GB-0011459
 87.5%;
                                                                                    AAS18869 standard; DNA; 18
                                                                                                                     (first entry)
                                          rcraggedercagicri
          Conservative
  Best Local Similarity
                                                                                                                                                                                                                               WO200185993-A2.
                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                               15-NOV-2001
                                                                                                                     12-MAR-2002
           14;
                                                                                                    AAS18869;
                                                                            AAS18869/c
          Matches
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The invention described a method of detecting variation in growth hormone (GH1), and therefore GH dysfunction in an individual. The method comprises comparing the nucleotide sequence of GH1 gene obtained from the test sample with a standard human GH1 gene sequence, in order to identify variation (GH1 variant). The method is useful in screening patients for treating irregularities or producing variant proteins for treating irregularities, and for the early detection and appropriate clinical management of familial GH deficiency. The GH1 variants are useful in chrappentic, diagnostic or detection method, particularly for determining binding defects and susceptibility to a disease such as diabetes, obesity or infection; for treating acromegaly or gigantism conditions associated with sodium and water retention, metabolic conditions associated with sodium and water retention, metabolic syndromes, mood and sleep disorders; diagnosing GH dysfunction and etectuling genetherapy or protein therapy. The GH1 variants are sepcially useful in gene therapy or protein therapy. The GH1 or GH variant may also be used in the preparation of a medicament, diagnostics composition or the used in the etiology of short stature; identifying of the mode of mutations in the etiology of short stature; identifying of the mode of metations in the structure and function of the GH molecule and development of arapid diagnostic tests for inherited GH deficiency. This sequence is the GH1 gene locus control region (LCR) specific primer, LCR25, used to amplify
Detecting growth hormone variants (GH1), useful in screening patients for growth hormone irregularities, comprises comparing the nucleotide sequence of a GH1 gene from a test sample with that of a standard sequence of the human GH1 -
                                                                                                                                                                                                                                                                                                                                            Claim 11; Page 77; 95pp; English.
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0

Gaps

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Indels

Score 12.8; DB 1; 1 Pred. No. 2.9e+02; 0; Mismatches 2;

Query Match 0.9%; Best Local Similarity 87.5%; Matches 14; Conservative

Sequence 18 BP; 4 A; 1 C; 11 G; 2 T; 0 other;

Length 18;

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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates, (b) a primer designed based on the chromosome marker or sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant corresponding to the marker is equence; (d) the multiwell plates containing the clones having said marker sequence; (d) the order of the marker is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell containination Nos. are mixed respectively in each wells of Inngitudinal and lateral directions; (f) the mixed clones are cultured and the cashltant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell cash detected from the amplified products; (h) the clones in the multiwell cash detected from the amplified products; (h) the clones in the multiwell cash detected from the amplified products; (h) the clones in the multiwell cash detected from the amplified products; (h) the clones in the multiwell cash detected from the amplified products; (h) the clones in the multiwell cash detected from the amplified products; (h) the clones in the multiwell cash detected from the multiwell cash detected from the 
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the LCR during sequence analysis to identify GH1 variants, described in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plates are specified from the detected result; and (1) the clones are
                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis;
                                                                                                                                                                                            ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention.
                                                                                                                                         Length 18;
                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human chromosome 1p36-35 PCR primer SEQ ID NO:188.
                                                                                                                                   . Match 0.9%; Score 12.8; DB 1; Local Similarity 87.5%; Pred. No. 2.9e+02;
                                                                                  Sequence 18 BP; 4 A; 7 C; 4 G; 3 T; 0 other;
                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; Page 8; 528pp; Japanese.
                                                                                                                                                                                                                                                      1255 TGAGGCCAGGTTGAGG 1270
                                                                                                                                                                                                                                                                                                                                                                                                                                                ABL43144 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-MAR-2001; 2001JP-0068285.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-MAR-2000; 2000JP-0066716.
                                                                                                                                                                                                                                                                                                         16 rickierckierrickie 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIKA ) RIKAGAKU KENKYUSHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                  14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Arraying genome clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genome; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-144136/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JP2001321190-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 654
                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                            ABL43144
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              88833
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Gregory J, Millar DS;

(UYWA-) UNIV WALES COLLEGE OF MEDICINE.

Procter AM,

Cooper DN,

WPI; 2002-089798/12.

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chromosome 1p36-35; chromosome 21q22.1; genetic analysis;
                                                                                 Human chromosome 1p36-35 PCR primer SEQ ID NO:1871.
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 6 A; 7 C; 3 G; 2 T; 0 other;
                                                                                                                                                                                                                              Claim 4; Page 41; 528pp; Japanese.
936 GGAGAAGAGGTGTGAG 951
            GGAGCAGGGGTGTGAG 17
                                                                                                                                                         12-MAR-2001; 2001JP-0068285
                                                                                                                                                                     10-MAR-2000; 2000JP-0066716
                                                                                                                                                                                  (RIKA ) RIKAGAKU KENKYUSHO.
(GENO-) GENOTEX YG.
                                            ABL44827 standard; DNA; 18
                                                                      (first entry)
                                                                                                       PCR primer; ss.
                                                                                                                                                                                                                   Arraying genome clones
                                                                                                                                                                                                      WPI; 2002-144136/19
                                                                                                                                JP2001321190-A.
                                                                                                                    Homo sapiens.
                                                                      11-APR-2002
                                                                                                                                             20-NOV-2001.
                                                         ABL44827;
                                                                                                       genome;
                                 RESULT 655
                                       a
  à
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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker of sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant containing the clones having said marker sequence; (d) the order of plates containing the clones having said marker sequence; (d) the order cof the maximum in the specified discrimination Nos. to array the multiwell cof the maximum in the specified discrimination Nos. to array the multiwell and lateral directions; (f) the mixed clones are cultured and the clones in the multiwell plates of the specified and lateral directions; (f) the mixed clones are cultured and the containing the amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABL43937 to ABL4332 represent profession of the contain the analysis of plates; and plates are specified for microarray is useful for gene analysis. ABL43937 to ABL4333 to ABL43634 enemity of analysis. specifically claimed for use in the present invention

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..
                          Gaps
                          ;
Length 18;
                            2; Indels
 0.9%; Score 12.8; DB 1;
87.5%; Pred. No. 2.9e+02;
tive 0; Mismatches 2;
                              14; Conservative
             Best Local Similarity
Matches 14; Conserv
      Query Match
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554 CAGGCATGCACACT 569
                               caegcarecacca 17
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ð g RESULT 656 ACA60576

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Human; mouse; ss; primer; gene 216; antiasthmatic; antiinflammatory; anorectic; chromosome 20p13-p12; single nucleotide polymorphism; SNP; gene therapy; respiratory disease; asthma; obesity; PCR; bronchial hyper-responsiveness; chronic obstructive pulmonary disease; adult respiratory distress syndrome; inflammatory bowel syndrome.
                                                                                                                                                                                                                                                                                                                                                                                              with
to 50
cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention describes a compound (I) of up to 50 nucleobases in length, which inhibits the expression of Cyclin D2. (I) is useful for inhibiting the expression of Cyclin D2 in cells or tissues in vitro. (I) is thus useful for treating disease associated with Cyclin D2 expression. (I) is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. This sequence represents human cyclin D2 inhibition associated oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                             Antisense inhibition of human cyclin D2 related oligonucleotide #13.
                                                                                                                                                                                                                                                                                                                                                                                           Novel antisense compound useful for treating diseases associated Cyclin D2 expression, comprises an oligonucleotide comprising up nucleobases in length, which inhibits expression of Cyclin D2 in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 0.9%; Score 12.8; DB 1; Length 18; Local Similarity 87.5%; Pred. No. 2.9e+02; Indels nes 14; Conservative 0; Mismatches 2; Indels
                                                                                                              Human; cyclin D2; diagnostic; therapeutic; prophylaxis; cyclin 2 inhibition; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human gene 216 polymorphism detection PCR primer #116.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 2 A; 0 C; 4 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 15; Column 45-46; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1145 TTTTTTTTGGAA 1160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABX75059 standard; DNA; 18 BP
 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rirrgrirrigga 18
                                                                                                                                                                                                                                                                                   01-AUG-2001; 2001US-0920760.
                                                                                                                                                                                                                                                     01-AUG-2001; 2001US-0920760.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
ACA60576 standard; DNA; 18
                                                           (first entry)
                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-361492/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   or tissues in vitro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200283077-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
                                                                                                                                                                                               JS6492173-B1.
                                                                                                                                                                   Homo sapiens.
                                                           11-JUN-2003
                                                                                                                                                                                                                           10-DEC-2002
                                                                                                                                                                                                                                                                                                                                               Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABX75059;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                ACA60576;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABX75059/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 657
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Keith T, Simon J,

us09904568-3.rng

Metz JG;

Page 322

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The present invention relates to novel nucleic acids and proteins for accolactate synthase (ALS) regulatory regions, alpha-tubulin promoter, polyketide synthase (PKS) promoter and fatty acid desaturase promoter from Thraustochytriales microorganisms. The nucleic acids of the invention are useful for transforming Thraustochytriales microorganisms or the foreign nucleic acids in a Thraustochytriales. The present sequence is a primer which is used to detect the presence of pUNBZEO11-2 sequences in Schizochytrium species. This sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a method for detecting a skin stimulation effect by using a test substance. The method comprises contecting a test substance with epithelial cells, and detecting the increase or decrease of calcium ion concentration in the cells. The method is useful as a
                                                                                                                                                            New nucleic acid molecule, useful for transforming Thraustochytriales microorganisms or the foreign nucleic acids in a Thraustochytriales
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Receptor-based method for detecting skin stimulation effect useful in screening analgesics, antiinflammatory agents and inhibitors on epidermal abnormality, comprising measuring changes in calcium ion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dermatological; analgesic; antiinflammatory; epithelial cell; skin; vallinoid receptor; VR1; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vallinoid receptor (VR1) related PCR primer 1 # SEQ ID 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%; Score 12.8; DB 1;
87.5%; Pred. No. 2.9e+02;
live 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 2 A; 6 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Denda S;
                                                                          Ramseier IM,
                                                                                                                                                                                                                                Example 4; Page 106; 112pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Examples; Page 8; 22pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Denda M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       993
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                                                                          Matthews TD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABZ56925 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SHIS ) SHISEIDO CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Inoue K, Fujiwara S,
                            (OMEG-) OMEGATECH INC
                                                                                                                      WPI; 2003-075541/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-140773/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2002103351-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               concentration
                                                                          Roessler PG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention relates to a novel isolated nucleic acid, gene 216, identified from human chromosome 20p13-p12. The invention also discloses regions of the 216 gene that contain single nuclectide polymorphisms (20PP's) which may be used as markers for disease susceptibility or severity. The nucleotides of the invention may have antiasthmatic, attinfiammatory or anorectic activities and may be used in gene therapy. The nucleic acids, antibodies or its fragments are useful for diagnosing, preventing or treating a discretz, such as respiratory diseases (e.g. asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease or adult respiratory distress syndrome, obstructive pulmonary disease or adult respiratory distress syndrome, obstructive nucleic acids are also useful for identifying increased susceptibility of a subject to the disorders mentioned. The nucleic acids can also useful for the recombinant production of disorder-associated templates for the recombinant production of disorder-associated templates for the recombinant production of disorder-associated peptides, for chromosome and gene mapping, or for tissue distribution studies. The present sequence represents a gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sb.
                                                                                                                                                                                                                                                                                                                                       New isolated gene 216 nucleic acids, useful for diagnosing, preventing or treating a disorder, such as asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease, obesity or inflammatory bowel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                to detect the presence of pTUBZE011-2 in Schizochytrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ALS; alpha-tubulin; polyketide synthase; PKS; primer; ss.
                                                                                                                                                                                                                                  Del Mastro RG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.9%; Score 12.8; DB 1; Length 18; 17.5%; Pred. No. 2.9e+02; ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               216 specific PCR primer used in the scope of the invention
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                                                                                                                                                                                                                                  Dupuis J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 2 A; 7 C; 3 G; 6 T; 0 other;
                                                                                                                                                                                                                                Van Eerdewegh P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 10; Page 156; 650pp; English.
                                                                                                                                                       (SCHE ) SCHERING CORP.
(GENO-) GENOME THERAPEUTICS CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP.
                                                                                                                                                                                                            Little RD, Van L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  725 AGCAGGGGGCCTGGCT 740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 Similarity 87.5%;
14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-APR-2002; 2002WO-US12040.
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                                                                                                              13-APR-2001; 2001WO-US12245.
                                                15-APR-2002; 2002WO-US12063
                                                                                           13-APR-2001; 2001US-0834597
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16 AGCAGAGGGCATGGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Acetolactate synthase;
fatty acid desaturase;
                                                                                                                                                                                                                                                                                                 WPI; 2003-092960/08.
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Best Local Similarity
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24-OCT-2002
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AAD50969;

RESULT 658

Matches

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Gaps

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Indels

Length 18;

us09904568-3.rng

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The present invention describes a method (M1) for regulating expression of a nucleic acid sequence in a primary cell comprising providing a nucleic acid binding polypeptide capable of binding to the nucleic acid sequence and contacting the nucleic acid binding polypeptide (I) and binding polypeptide (I) and in a nucleic acid sequence in the primary cell. Also described: (I) a nucleic acid sequence in a primary cell; (2) a primary cell (II) comprising an exogenous nucleic acid binding polypeptide; (2) a primary cell (II) comprising an exogenous nucleic acid binding polypeptide or the primary cell and a carrier or diluent; (4) treating (M2) or preventing a disease, or (5) expression (II) comprising maxy cell. (I) has cytostatic and nephrotropic polypeptide in a primary cell. (I) has cytostatic and nephrotropic treating or preventing a disease e.g., kidney failure or cancer. The present sequence represents an oligonocleotide which is useful the
                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Regulating expression of a nucleic acid sequence in a primary cell for treating or preventing a disease e.g., cancer, comprises contacting the nucleic acid binding polypeptide with the nucleic acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gene regulation; expression; nucleic acid binding protein; cytostatic; nephrotropic; gene therapy; kidney failure; cancer; ss.
dermatological agent in the screening of analgesics, antiinflammatory agents and inhibitors on epidermal abnormality. The current sequence represents a rat vallinoid receptor (VRI) related PCR primer sequence
                                                                                                                                                                               Gaps
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                                                                                                                                   . Match 0.9%; Score 12.8; DB 1; Length 18; Local Similarity 87.5%; Pred. No. 2.9e+02; les 14; Conservative 0; Mismatches 2; Indels
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                                                                                               Sequence 18 BP; 5 A; 7 C; 5 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     exemplification of the present invention.
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                                                             used in an example from the invention.
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                                                                                                                                                                                                                546 CCTGCTGGCAGGCATG 561
                                                                                                                                                                                                                                          18 CCTGCTGGGTGGCATG 3
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                       agents and
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Matches
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABR099989, ABR00010-ABH99989 and ABL001010-ABH99989 and NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fip.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            oet or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single nucleotide polymorphisms and cytosine methylation status
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                                                                                                                                                                                                                    Oligonucleotide SEQ ID NO 187741 for detecting SNP TSC0046249.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID 187741; 29pp + Sequence Listing; German.
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298 TCTGCTGTGGGGGCTG 313
                                 1 rcregesreesecre 16
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ABF87744 standard; DNA; 13
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Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (EPIG-) EPIGENOMICS AG
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                                                                                                  ABF87744/
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AC ABF8
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0.9%; Score 12.8; DB 1; Length 18; larity 87.5%; Pred. No. 2.9e+02; Conservative 0; Mismatches 2; Indels

Query Match Best Local Similarity Matches 14; Conserv

06-APR-2001; 2001WO-IB00713

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                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                               Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single nucleotide polymorphisms and cytosine
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              Oligonucleotide SEQ ID NO 187742 for detecting SNP TSC0046249.
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0.9%; Score 12.6; DB 1; Length 13;
Best Local Similarity 92.3%; Pred. No. 2.1e+02;
Matches 12; Conservative 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                               Berlin K;
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                                                                                                                                                                                                                                                                                           (EPIG-) EPIGENOMICS
                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-657177/75
                                                                                                                                                                                                                                                                                                                                                                                                                                 methylation status
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                                                                                                                      Homo sapiens
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosline methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal respiratory, oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF0010-ABH99999 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this parent did not form part of the printed specification, the unvention. Appendigued the oligoners are also used for detains the oligomers are also used for detains described in the invention.
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                                                                                                                                                                                                                                                                                                                     Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single nucleotide polymorphisms and cytosine methylation status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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92.3%; Pred. No. 2.1e+02;
ve 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID 193455; 29pp + Sequence Listing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13 BP; 2 A; 0 C; 2 G; 8 T; 1 other;
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                                                                                                                                                                                              Berlin K;
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                                                                                                                                                                                              Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Piepenbrock C,
                                                                                                                          (EPIG-) EPIGENOMICS AG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200177384-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABF93459;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Olek A,
                                                                                                                                                                                              olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 66
ABF93459/
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, clentral nervous system, cardiovascular and metabolic disorders. The ABCOO10-ABC99989, ABF0010-ABE9989, ABF0010-ABE9989, ABF0010-ABE9989, ABF0010-ABE9989, and NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                    Claim 1; SBQ ID 193456; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                              ftp.wipo.int/pub/published_pct_sequences
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Sequence 13 BP; 8 A; 2 C; 0 G; 2 T; 1 other;

. 0 0.9%; Score 12.6; DB 1; Length 13; 22.3%; Pred. No. 2.1e+02; Iv Mismatches 0; Indels Query Match 0.9%; Best Local Similarity 92.3%; Matches 12; Conservative

1104 TTATGTAGTTTTC 1116 

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999 RESULT

ABH10200 standard; DNA; 13

ABH10200;

BP.

(first entry) 22-FEB-2002

Oligonucleotide SEQ ID NO 210177 for detecting SNP TSC0051320.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens,

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB00713

2000DE-1019173 07-APR-2000;

(EPIG-) EPIGENOMICS AG.

Berlin K; Olek A, Piepenbrock C, 

WPI; 2001-657177/75.

typing, is cytosine Set of oligonucleotides, useful for diagnosis and cell designed to detect single nucleotide polymorphisms and methylation status

Claim 1; SEQ ID 210177; 29pp + Sequence Listing; German.

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The ABC00010-ABC99989, ABF00010-ABP9989, ABH00010-ABH99989 and ABC00010-ABB9989, ABF00010-ABH99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed

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                                                                                          Gaps
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specification, but was obtained in electronic format from WIPO
                                                                                          .
0
                                                                  Length 13;
                                                                                        Indels
                                                                                      0
                                                              Score 12.6; DB 1;
Pred. No. 2.1e+02;
                                                      0.9%; Scor.
92.3%; Pred. No. 2...
                                    Sequence 13 BP; 4 A; 0 C; 3 G; 5 T; 1 other;
              ftp.wipo.int/pub/published_pct_sequences
                                                                                     12; Conservative
                                                            Query Match
Best Local Similarity
Matches 12; Conserv
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ABH10201 standard; DNA; 13

BP.

ABH10201;

22-FEB-2002 (first entry)

0;

Gaps

Oligonucleotide SEQ ID NO 210178 for detecting SNP TSC0051320.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2

18-OCT-2001.

06-APR-2001; 2001WO-IB00713.

07-APR-2000; 2000DE-1019173.

(EPIG-) EPIGENOMICS AG

Berlin K; Piepenbrock C, olek A,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single nucleotide polymorphisms and cytosine methylation status

Claim 1; SEQ ID 210178; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABL00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed appendication, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published\_pct\_sequences. RESULT 66
RESULT 66
REH 10201/
ID ABH1
XX ABH1
XX ABH1
XX SNP;
XX SNP;
XW Cent
XX NW Cent
XX Cent
XX

Sequence 13 BP; 5 A; 3 C; 0 G; 4 T; 1 other;

Gaps ; 0 0.9%; Score 12.6; DB 1; Length 13; 32.3%; Pred. No. 2.1e+02; Ive 1; Mismatches 0; Indels 92.3%; Best\_Local Similarity 92.3 Matches 12; Conservative Query Match

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1100 GTAATTATGTAGT 1112

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Hodgkin's disease; HD; acid phosphatase deficiency; noveb polymorphic site; ACP2 hablotype; ACP2 genotype; polymorphism; transgenic animal; primer; probe; primer-extension oligonucleotide; SNP; single nucleotide polymorphism.

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The invention comprises the amino acid and coding sequence of the human interleukin 12A (II-12A) protein. Specifically the invention relates to the identification of polymorphisms within the human (II-12A) gene sequence. The polymorphisms identified in the human II-12A gene sequence are useful in studying the expression and function of II-12A gene sequence screening drugs for the treatment of disorders such as AIDS, malaria, tuberculosis and cancer. The II-12A polymorphisms may be used to haplotype and genotype the II-12A gene of an individual. The II-12A DNA sequences of the invention can be used to create transgenic animals for setudying expression of the II-12A isogenes in vivo. The present DNA sequence represents a human interleukin 12A (II-12A) gene allele specific oligonucleotide primer.
                                                                                                                                      Human interleukin 12A (IL-12A) allele specific oligonucleotide primer 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New interleukin 12A (IL-12A) gene polymorphic variants, for studying the expression and function of IL-12A and screening candidate drugs for
                                                                                                                                                                          Human; primer; interleukin 12A; IL-12A; drug screening; AIDS; malaria
tuberculosis; cancer; haplotyping; genotyping; transgenic animal; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Koshy B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kliem SE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Choi JY, Gilson CR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 BP; 3 A; 2 C; 5 G; 4 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 15; Page 13; 72pp; English.
                                   AAL44238 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                             (GENA-) GENAISSANCE PHARM INC.
                                                                                                                                                                                                                                                                                                                                         05-OCT-2001; 2001WO-US31656.
                                                                                                                                                                                                                                                                                                                                                                           06-OCT-2000; 2000US-238693P.
                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cappola G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              treating AIDS and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-315865/35.
                                                                                                                                                                                                                                                                    WO200229115-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Armstrong B,
                                                                                                                                                                                                                                  Homo sapiens
                                                                                                        08-NOV-2002
                                                                                                                                                                                                                                                                                                        11-APR-2002
                                                                       AAL44238;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Parks KE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
RESULT 667
                    AAL44238
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for

Novel genetic variants of acid phosphatase 2, lysosomal polypeptide gene useful in studying expression and function of the protein, and screening drugs to treat diseases e.g. Hodgkin's disease

Tanguay DA;

Меввег С,

Kliem SE,

WPI; 2002-154563/20.

(GENA-) GENAISSANCE PHARM INC.

07-JUN-2001; 2001WO-US18457. 07-JUN-2000; 2000US-210047P.

WO200194362-A2 Homo sapiens

13-DEC-2001,

Claim 17; Page 14; 109pp; English.

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The invention comprises the human lysosomal acid phosphatase 2 (ACP2) nucleic acid and protein sequences. Specifically, the invention relates to the discovery of 22 novel polymorphic sites within the APC2 gene. The invention also comprises methods for haplotyping and genotyping the ACP2 gene in an individual. The ACP2 gene (located on chromosome 11) encodes a CP sposomal-specific enzyme that catalyses the hydrolysis of critically important in the treatment of Hodgkin's protein are pharmaceutically important in the treatment of Hodgkin's corporation are pharmaceutically important in the treatment of Hodgkin's corporation of the invention are useful in haplotyping the ACP2 gene. Copymorphisms of the invention are useful in haplotyping the ACP2 gene. ACP2 haplotyping is useful in validating ACP2 as a target (and designing disease and acid phosphatase deficiency.) The ACP2 gene polymorphisms of the invention are useful in the production of a transpencin acids of the invention are useful in the production of a transpencin and hich expresses ACP2 protein. The ACP2 protein and nucleic acids of the invention are useful in the production of allele-specific oligonucleotides designed to genotype each of the ACP2 protein. The ACP2 uncleic acids ABL36329-ABL3634 represent claimed ACP2 allele-specific process. Nucleic acids ABL36321-CCC ABL3634 represent claimed ACP2 allele-specific process. Nucleic acids ABL36321-CCC ABL36354 represent claimed ACP2 allele-specific process. Nucleic acids ABL36321-CCC ABL36354 represent claimed ACP2 allele-specific process.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.9%; Score 12.6; DB 1;
92.3%; Pred. No. 2.6e+02;
iive 1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15 BP; 2 A; 8 C; 3 G; 1 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      202
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 GCCGCCACCRG 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  190 GCCGCCCACCCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAD44145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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ZXEXEXEX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human lysosomal acid phosphatase 2 (ACP2) allele-specific PCR primer 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ó;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12.6; DB 1;
Pred. No. 2.6e+02;
1; Mismatches 0;
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. 98; 92.3%;

744 GCATGTTGCTGAC 756

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Conservative

Local Similarity

12;

Matches

GCATGTTGCTGAY 14

ö

Gaps

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Length 15; Indels Oligo-dT PCR primer #5 used to illustrate the method of the invention.

Sequential consensus region-directed amplification; gene expression;

Human; ss; lysosomal acid phosphatase 2; ACP2; gene; chromosome 11; lysosome-specific enzyme; orthophosphoric monoester hydrolysis;

ABL36325 standard; DNA; 15 BP.

RESULT 668

ABL36325 ID ABL

(first entry)

22-APR-2002

MAXEXEX MAXEX MAXE

ABL36325;

WPI; 1999-430152/36.

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The invention relates to a method of sequential consensus region-directed amplification for sorting a mixture of DNAs into 2 or more subsets or distinguishing gene expression patterns in 2 samples. The methods, kits and oligonucleotides are useful for sorting a mixture of DNAs into 2 or more subsets or distinguishing gene expression patterns in 2 samples e.g. for disease diagnosis and gene analysis. The present sequence is oligo dT PCR primer used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                        Sequential consensus region-directed amplification for sorting mixture of DNAs into 2 or more subsets or distinguishing gene expression patterns in 2 samples, useful for disease diagnosis and gene analysis
disease diagnosis; gene analysis; human; matrix metalloproteinase;
                                                                                                                                                                                                                         (UYVI-) UNIV VIRGINIA COMMONWEALTH INTELLECTUAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 16 BP; 0 A; 1 C; 0 G; 14 T; 1 other;
                                                                                                                                                                                                                                                             W, Gillies G;
                                                                                                                                                                                                                                                                                                                                                                                             Example, Fig 1C, 19pp, English.
                                                                                                                                                                                         97US-108152P
                                                                                                                                                                                                                                                           Broaddus
                                                                                                                                                                                                                                                                                           WPI; 2002-412824/44.
                   PCR; primer; ss.
                                                   Unidentified
                                                                                                                                                                                       03-OCT-1997;
                                                                                   US6277571-B1
                                                                                                                                                     30-SEP-1998;
                                                                                                                                                                                                                                                       Fillmore H,
                                                                                                                     21-AUG-2001
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Gaps
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0
ch 0.9%; Score 12.6; DB 1; Length 16; l Similarity 86.7%; Pred. No. 2.8e+02; 13; Conservative 1; Mismatches 1; Indels
                                                                       1142 CCTTTTTTTTTT 1156
                                                                                                              1 CVTTTTTTTTTTT
 Query Match
Best Local Similarity
Matches 13; Conserv
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Reverse transcription primer Pl
                BP.
               AAX87332 standard; DNA; 18
                            (first entry)
                           27-SEP-1999
                     AAX87332;
         RESULT 670
d
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SAG gene; sensitive to apoptosis; mouse; cancer; tumour; neurodegenerative disease; muscular dystrophy; wound healing; vulnerary; therapy; PCR; primer; ss.

WO9932514-A2 Synthetic

01-JUL-1999

15-DEC-1998;

98WO-US26705

98US-0099840. 19-DEC-1997; 11-SEP-1998;

LAMBERT CO. (WARN ) WARNER

Sun Y;

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Nootropic; neuroprotective; antiparkinsonian; anticonvulsant; analgesic; tranquiliser; antidiabetic; ophthalmological; neurodegenerative disorder; neublastin; ischemic neuronal damage; traumatic brain injury; diabetes; peripheral neuropathy; neuropathic pain; Alzheimer's disease; glaucoma; Huntington's disease; Parkinson's disease; amyotrophic lateral sclerosis; memory impairment; renal disease; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a truncated neublastin polypeptide comprising an
                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New truncated neublastin polypeptides lacking one or more amino-terminal amino acids of a mature neublastin polypeptide useful for treating neurodegenerative disorders, e.g. peripheral neuropathy,
                                                                                                             This primer was used for reverse transcription of RNA isolated from mouse tumour lines L-RT101 (epidermal tumour cell line) and H-TX (spontaneously transformed liver line). It was also used as the reverse primer in PCR amplification of the resulting cDNA. Primers P1 and P2 (see AAX87333) reproducibly detected differential expression of a gene between 1,10-phenanthroline (OP)-treated and OP-nontreated L-RT101 and H-Tx cells. An OP-inducible clone was oorresponding to the mouse sensitive to apoptosis gene (SAG): SAG is a redox-sensitive, haem-binding protein domain that promotes cell growth, protects cells from apoptosis, scavenges oxygen radicals and can be used for the reversion of a tumour phenotype.
                               SAG: Sensitive to Apoptosis Gene and related proteins, useful for promoting cell growth and protecting cells against apoptosis
                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                      Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                  0.9%; Score 12.6; DB 1; 86.7%; Pred. No. 3.2e+02; ive 1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 2 A; 1 C; 1 G; 13 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rossomando A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neublastin DNA related PCR primer.
                                                                                  Example 1; Page 14; 84pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure, Fig 8; 138pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neuropathic pain, brain injury
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       1143 CTTTTTTTTTTG 1157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-MAR-2001; 2001US-0804615
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-MAR-2002; 2002WO-EP02691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABT11916 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                     13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sah DWY, Johansen TE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-713515/77.
                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (BIOJ ) BIOGEN INC
(NSGE-) NS GENE AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200272826-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 671
ABT11916/c
                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
# X E E X & X & C C C C C C C C X & S
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a mature neublastin polypeptide. The polypeptides and nucleic acids are useful for treating neurodegenerative disorders such as ischemic neuronal damage, traumatic brain injury, peripheral neuropathy, neuropathic pain, Alzheimer's disease, Huntington's disease, Parkinson's disease, amyotrophic lateral sclerosis, memory impairment, diabetes, renal diseases, or glaucoma by moderating metabolism, growth, differentiation or survival of a nerve or neuronal cell. This polymucleotide sequence is a neublastin PCR primer of the invention. acid terminus that lacks one or more amino-terminal amino acids of 8X88888888XX

Sequence 18 BP; 1 A; 6 C; 9 G; 2 T; 0 other;

Gaps . 0 Length 18; 1; Indels 0.9%; Score 12.4; DB 1; 92.9%; Pred. No. 3.5e+02; Mismatches ; 806 CCCTGCAGCCGAGC 819 ß Conservative CCCTGCAGCCGGGC Query Match Best Local Similarity 13; 18 Matches

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RESULT 672

ABK94277 standard; DNA; 21 ABK94277; Endothelin converting enzyme 1 (ECE-1) SNP detection primer #65

(first entry)

27-AUG-2002

Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; anadiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; single nucleotide polymorphism; PCR; primer; ss.

Synthetic.

WO200224747-A2.

28-MAR-2002

31-AUG-2001; 2001WO-EP10087

19-SEP-2000; 2000EP-0120123

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Brinkmann U, Hoffmeyer S;

WPI; 2002-435060/46.

Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful for signaling system rating the disease 

Claim 1; Page 63; 190pp; English.

The invention describes a polymucleotide (I) of the endothelin (EDN)/endothelin converting enzyme (ECB)/receptors of EDN and ECB (EDNR) signaling system which is associated with a cardiovascular disease. (I), the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing (I) or (II) is useful for producing cells capable of expressing a molecular variant polypeptide which is associated with a cardiovascular disease. (II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing a molecular variant gene comprising (I) is useful for identifying and obtaining a pro-drug or drug capable of modulating the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product, or for identifying and obtaining an inhibitor of the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE

ö signaling system or its gene product. The isolated proteins and polynucleotides encoding them are useful for preparation of a pharmacutical composition for treating a cardiovascular disease such as coronary heart disease, hypertension, atherosclerosis, or related to abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial hypercholesterolaemia. The gene or a polynucleotide fragment of the EDN/ECE/EDNR signaling system are useful as forensic markers, for creating a transgenic animal and in creation of a solid support comprising polynucleotides, genes, vectors, polypeptides, antibodies or host cells of the invention. This sequence represents a PCR primer used to identify single nucleotide polymorphisms in DNA encoding cardiovascular regulator proteins of the EDN/EC/EDNR signaling pathway. Gaps ., Length 21; 1; Indels Score 12.4; DB 1; Pred. No. 4.1e+02; 0; Mismatches 1; Sequence 21 BP; 6 A; 3 C; 11 G; 1 T; 0 other; .98; Query Match 

92.9%; 815 Local Similarity 92.9 les 13; Conservative 802 CGCTCCCTGCAGCC Matches ð

crerecerdedec s

8

RESULT 673 ABK94278

ABK94278 standard; DNA; 21

BP.

ABK94278;

(first entry) 27-AUG-2002

Endothelin converting enzyme 1 (ECE-1) SNP detection primer #66.

Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; BDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclenosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; single nucleotide polymorphism; PCR; primer; ss.

Synthetic

WO200224747-A2

28-MAR-2002.

31-AUG-2001; 2001WO-EP10087. 

19-SEP-2000; 2000EP-0120123

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

s, Hoffmeyer Brinkmann U,

WPI; 2002-435060/46.

for Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful disease the treating

Claim 1; Page 63; 190pp; English.

The invention describes a polynucleotide (I) of the endothelin (EDN) fendothelin converting enzyme (ECE)/receptors of EDN and ECE (EDNR) signaling system which is associated with a cardiovascular disease. (I), the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing (I) or (II) is useful for producing cells capable of expressing a molecular variant polypeptide which is associated with a cardiovascular disease. (II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing a molecular variant gene comprising (I) is useful for identifying and obtaining a pro-drug or drug capable of modulating the activity of a

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molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system
or its gene product, or for identifying and obtaining an inhibitor of
the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE
signaling system or its gene product. The isolated proteins and
polymucleotides encoding them are useful for preparation of a
polymucleotides encoding them are useful for preparation of a
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polymucleotides comparation of a series of the series of polymucleotide fragment of the
coronary heart disease, hypertension, atherosclerosis, or related to
abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial
hypercholesterooleaning system are useful as forensic markers, for
creating a transgenic animal and in creation of a solid support
creating a transgenic animal and in creation of a solid support
comprising polymucleotides, genes, vectors, polypeptides, antibodies or
comprising to the invention. This sequence represents a PCR primer used
to identify single nucleotide polymorphisms in DNA encoding
cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway.
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2.9%; Pred. No. 4.1e+02;
ve 0; Mismatches 1;
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Human flt1 VEGF receptor hairpin ribozyme substrate #62. BP. AAX70294 standard; RNA; 18 (first entry) 28-JUL-1999 AAX70294; RESULT 674 AAX70294/c 

Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.

Homo sapiens.

WO9715662-A2

01-MAY-1997

96WO-US17480 25-OCT-1996;

96US-0584040. 95US-0005974. 11-JAN-1996; 26-OCT-1995;

RIBOZYME PHARM INC. (CHIR ) CHIRON CORP. (RIBO-) RIBOZYME PHZ McSwiggen J, Pavco P, Stinchcomb D; Escobedo J,

WPI; 1997-259017/23.

Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient

Claim 4; Page 94; 218pp; English

The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VBGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (fit-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (fik-1) (e.g. tumour

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ligand dependent transcriptional factor; oestrogen receptor; ER; glucocorticoid receptor protein; GR; mineralocorticoid receptor protein; MR; peroxisome proliferator-activated receptor protein; PPAR; progreterone receptor protein; PR; pregname X receptor protein; PXR; thyroid hormone receptor protein; TR; vitamin D receptor protein; VDR; transactivation; ERalpha; breast cancer; PCR primer; probe; ss.
angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ligand dependent transcriptional factors, nucleic acids encoding them and cells comprising them and a specified reporter gene, useful for screening agents for the treatment of breast cancer -
                                                                                                                                                                     Gaps
                                                                                                                                                                     ;
0
                                                                                                                                 Length 18;
                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                            Human oestrogen receptor alpha probe oligonucleotide 31
                                                                                                                           Score 12.2; DB 1;
Pred. No. 3.8e+02;
0; Mismatches 3;
                                                                                         Sequence 18 BP; 2 A; 4 C; 6 G; 6 U; 0 other;
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                                                                                                                                                                                                      1027 CAAAGIGCAGCIGACIC 1043
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07-JUL-2000; 2000JP-02205011.
21-JUL-2000; 2000JP-02226508.
02-AUG-2000; 2000JP-0234653.
03-AUG-2000; 2000JP-0235460.
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                                                                                                                                                82.4%;
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                                                                                                                                                                                                                                                                                                                               AAH57086 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000JP-0235463
                                                                                                                                                                                                                                      CAAAAAGCAGCTGGCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ohe N, Satoh H;
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                                                                                                                                              Local Similarity 82.4
nes 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-367866/38.
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transactivation of a reporter gene by a test ERalpha, for screening mutant ligand dependent transcriptional factors, for evaluating an activity for transactivation of a reporter gene by a test ERalpha and/or for screening a compound useful for treating a disorder of a mutant
                                                                                                        especially breast cancer.
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Length 20; Indels 0.9%; Score 12.2; DB 1; 12.4%; Pred. No. 4.2e+02; ve 0; Mismatches 3; Sequence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 other; 916 GCAGGCCTGACCCTGCA 19 82.4%; 960 GCAGGACTGACCCCTCA Best Local Similarity 82.4 Matches 14; Conservative m Query Match

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Gaps

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AAA11329/ RESULT

AAA11329 standard; DNA; 20 BP (first entry) 08-NOV-2000 AAA11329 

Human TRPC7 gene exon 23/intron 23 junction.

Transmembrane protein, TRPC7; brain; transient receptor potential; TRP; calcium channel function; human; gene therapy; periodic psychosis; mutation; ss.

Homo sapiens

Socation/Qualifiers 11..20 /\*tag= b /number= 23 /\*tag= a
/number= 23 1..10 /\*tag= intron exon

WO200029571-A1

11-NOV-1999; 25-MAY-2000

99WO-JP06289 98JP-0321200 12-NOV-1998;

(EIKE ) EIKEN KAGAKU KK.

Nagamine K; Shimizu N,

WPI; 2000-387784/33.

Nucleic acids encoding transmembrane protein TRPC7 expressed in brain and homologous to transient receptor potential protein useful in the treatment of associated diseases such as periodic psychosis

Example 7; Page 39; 77pp; Japanese.

The invention relates to the isolation of a nucleic acid (AAA11284) coding for a transmembrane protein TRPC7 (AAY92944) which is expressed in brain and is homologous to transient receptor potential (TRP) protein. This suggests that the TRPC7 protein may have a calcium channel function. The genomic sequence has been shown to contain 31 introns. This sequence represents an exon/intron junction from the genomic TRPC7 sequence. The DNA and protein can be used in the diagnosis and treatment of disorders associated with TRPC7, especially the screening, monitoring and treatment (by gene therapy) of periodic psychosis, which appears to be associated with mutations in the TRPC7 gene.

New antisense compound targeted to nucleic acid encoding mitogen-activated protein kinase 4, useful for treating immunologic disorder, inflammatory disorder or cancer

Claim 3; Page 93; 132pp; English.

Sequence 20 BP; 4 A; 3 C; 11 G; 2 T; 0 other;

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                                                                                                                                                                      Human; MEKK4 modulation; mitogen-activated protein kinase kinase 4; MTK1;
MAP3744, MAP three kinase 1; MAP/EKK kinase kinase 4; MAPKK74; cytostatic;
prophylaxis; immunological; Myperproliferative disorder; cancer; therapy;
antisense; inflammatory; phosphorothioate backbone; ss.
                    Gaps
                    0;
 Score 12; DB 1; Length 20;
Pred. No. 4.5e+02;
0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
/note= "2'-methoxyethyl nucleotides"
                                                                                                                                                     Human MEKK4 antisense oligonucleotide, ISIS #123152.
                                                                                                                                                                                                                                                                     /mod_base= OTHER
/note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wyatt JR;
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/note== "2'-methoxyethyl
                                                                                                                                                                                                                                           Location/Qualifiers
                                     TGGGAGCCAGCGGCCCTCC 706
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                                                  20 TCGGACCCACCTGCTCCTC 1
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                                                                                                  BP.
0.9%;
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/mod_base= r
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                                                                                                 AAD37217 standard; DNA; 20
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/mod_base=
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Query Match
Best Local Similarity 75.0°
Matches 15; Conservative
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                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                           Synthetic.
                                                                                                                   AAD37217;
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                                                                                 RESULT 677
                                                                                          AAD372
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331

Page

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The present invention relates to antisense compounds, compositions and methods for modulating the expression of MEKK4 (also referred as mitogenactivated protein kinase kinase 4; MAPZKK4; MAP Three kinase 1; MAPZEK kinase kinase 4; MAPZKK4; MTI. The antisense cligos are useful for inhibiting the expression of MEKK4 in cells or tissues. They are also with MEKK4 such as immunological, inflammatory, hyperproliferative disorder or cancer. Sequences of the invention are also useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. They are also useful in antisense therapy. The present sequence is an antisense oligomoclade targetted to human MEKK4 DNA. This sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 2 A; 4 C; 10 G; 4 T; 0 other;
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0.9%; Score 12; DB 1; Length 20;
75.0%; Pred. No. 4.5e+02;
ive 0; Mismatches 5; Indels
                                     0; Mismatches
                                                                  659 TGGTCGGGGACTTGGCCAGC 678
                                                                                                1 rédrégaddadereder 20
                 75.0%;
                                      Conservative
Query Match
Best Local Similarity
Matches 15; Conserv
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0

Gaps

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RESULT 678 AAL47730/c

AAL47730 standard; DNA; 21 

AAL47730;

BP.

(first entry) 18-SEP-2002

Ras gene PCR primer SEQ ID NO: 26.

K-ras; N-ras; H-ras; ras; oncogene; mutation detection; PCR; primer; probe; restriction mediated selection PCR; cancer; ss.

Unidentified.

WO200229005-A2.

11-APR-2002

02-OCT-2000; 2000US-237416P.

02-OCT-2001; 2001WO-US42422.

(ORTH ) ORTHO CLINICAL DIAGNOSTICS INC.

Fuery CJ; Todd AV, Belly RT,

WPI; 2002-479599/51

Amplifying and determining mutant sequences in DNA sample using thermostable restriction enzyme so that during thermocycling mutant sequences are enriched while wild-type sequences and/or primer induced sites are cleaved

R; Liu X;

Kekuda

AM, Malyankar UM, Kekuda Burgess CE, Zerhusen BD, nithson G, Li L, Ji W;

Shimkets RA, Burgess CI Boldog FL, Smithson G,

Vernet CAM,

Patturajan M,

Gangolli EA,

Anderson D, Casman SJ,

Spytek KA, Stone DJ,

WPI; 2002-590744/63

or

Novel isolated NOVX polypeptide useful for treating cardiomyopathy, atherosclerosis, metabolic disorders, diabetes, obesity, infectious disease, anorexia, neurodegenerative disorders, Alzheimer's disease

Example 1; Page 198; 318pp; English.

cancer

Claim 1; Page 74; 116pp; English.

The present invention relates to a method of amplifying and determining target mutant Ras sequences in a DNA sample, involving the use of a thermostable restriction enzyme and primers shown in AAL47705-AAL47771. The method used is designated restriction mediated selection polymerase chain reaction (REMS-PCR). The method can be used to detect H-ras, K-ras and N-ras mutations, which may lead to cancer. The present sequence is a PCR primer useful in the method of the invention.

Sequence 21 BP; 3 A; 9 C; 6 G; 3 T; 0 other;

The present invention relates to new NOVX polypeptides. The invention is useful for treating or preventing a NOVX-associated disorder such as cardicomyopathy or atherosclerosis, where the disorder is related to cell signal processing and metabolic pathway modulation in a subject, preferably human. The invention is also useful for treating metabolic disorders (e.g. obesity), diabetes, inflectious disease, neurodegenerative disorders (e.g. Alzheimer's disease, Parkinson's disease), immune

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0.9%; Score 12; DB 1; Length 21; 75.0%; Pred. No. 4.7e+02;
Query Match
Best Local Similarity
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                                                                                                                                                                                                                                             cell signal processing, metabolic pathway modulation, metabolic disorder, besity, diabetes; infectious disease, neurodegenerative disorder, anne, Alzheimer's disease; Parkinson's disease; immune disorder; acnoe; Alzheimer's disease; Parkinson's disease; immune disorder; cancer; haematopoietic disorder; circuitis; pancratitis; learning defect; memory defect; infertility; congenital heart defect; hair growth; pigmentation disorder; emborcine disorder; respiratory disease; health; gastro-intestinal disease; reproductive; neurological disease; neptrological disorder; endocrine disorder; alerge; inflammation; nephrological disorder; procession is disorder; and procession is disorder; in adiporter complement related CIQ tumour necrosis factor; out at first; beta adrenergic receptor kinase; EphAo(-Mek-2; glucose transporter; type la membrane-sushi domain containing; PCR; primer; ss.
                                                                                                                                                                                                                                   Human; NOVX; NOVX-associated disorder; cardiomyopathy; atherosclerosis;
  Gaps
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0
                                                                                                                                                                                                         Human adipocyte Clq Tumour Necrosis Factor-like PCR primer 1.
  Indels
  5.
  Mismatches
                           529 GAGGAGCAGCTGGGTGCCCT 548
                                                      20 GAGGGCCGCTGGGTCACAT 1
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02-MAY-2001; 2001US-288153P.
29-MAY-2001; 2001US-294075P.
24-JUL-2001; 2001US-307506P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-AUG-2001; 2001US-311590P.
10-AUG-2001; 2001US-311613P.
29-AUG-2001; 2001US-315617P.
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                                                                                                                        ABS52111 standard; DNA; 18
                                                                                                                                                                               (first entry)
  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CURA-) CURAGEN CORP.
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0

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4

Mismatches

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14; Conservative

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The molecules of

Gaps

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disorders, haematopoietic disorders and various cancers. The molecules of the invention are also useful for treating or preventing dirthösis, pancreatinis, learning and memory defects, infertility, congenital heart defects, acme, hair growth, pigmentation disorders, endocrine disorders, respiratory disease, gastro-intestinal diseases, reproductive, health, neurological diseases, bone marrow transplantation, endocrine diseases, allergy and inflammation, nephrological disorders system disorders, neuropsychiatric disorders and age-related disorders.

The present nucleic acid sequence represents a PCR primer that was used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A new isoform of myotonic dystrophy protein kinase includes a sequence encoded by exon 16 of the gene and is useful to detect presence or risk of myotonic dystrophy, myocardial infarction or a condition associated with muscle damage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Myotonic dystrophy, DM; protein kinase, DMPK; myocardial infarction; muscle damage; dysfunction; reverse transcriptase PCR; RT-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Myotonic dystrophy protein kinase (DMPK) isoform, primer 57.
                                                                                                                                                                                                0.9%; Score 11.8; DB 1; Length 18; 86.7%; Pred. No. 4.4e+02;
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                                                                                                                                                                     Sequence 18 BP; 3 A; 8 C; 5 G; 2 T; 0 other;
                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                       in the methods of the invention.
                                                                                                                                                                                                                                                                                                                                                                            ABK85826 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                  1297 CAGCCTGGCCCCATG 1311
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tiscornia G;
                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                    Ouery Match 0.9
Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MAHA/) MAHADEVAN M S. (TISC/) TISCORNIA G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-507644/54.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mahadevan MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                         24-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                          ABK85826;
                                                                                                                                                                                                                                                                                                                                             RESULT 680
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is a primer for the PCR amplification of exon 1 from the human interleukin 9 (hIL-9), also known as asthma associated factor 1, gene. hIL-9 plays a role in arboic allergy, asthma and related disorders, e.g. bronchial hyperesponsiveness (BHR), rhinitis, urticaria, allergic inflammatory disorders of the bowel and various forms of eczema. A naturally occurring polymorphism has been identified at position 117 demonstrate, individuals homozygous for Met at position 117 demonstrate, e.g. a lack of asthma and low serum immunoglobulin E (IgE) levels, while Thr/Thr homozygotes and Thr/Met heterozygotes are susceptible to
                                                                                                                                                                                                                                                  Human; interleukin 9; asthma associated factor 1; IL-9; primer; atopic allergy; asthma; bronchial hyperesponsiveness; BHR; eczema; rhinitis; urticaria; allergic inflammation; bowel; amplification; polymorphism; polymerase chain reaction; PCR; exon 1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human interleukin-9 variant with Met at position 117 - useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%; Score 11.6; DB 1; Length 18; 77.8%; Pred. No. 4.8e+02; ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                     Lower primer for exon 1 of human interleukin 9 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence 18 BP; 2 A; 11 C; 2 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Prasad KU;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        treating atopic allergy, esp. asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 42; 142pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nicholas N,
1275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    819
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                                                                                                                    AAT87896 standard; DNA; 18 BP.
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 1258 GGCCAGGTTGAGGCCCTT
                                  1 eggcágarggagggccrr
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                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lee MW, Levitt RC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1997-179278/16
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Local Sim-
14;
                                                                                                                                                                                      12-JAN-1998
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                                                                                                                                                                                                                                                                                                                                               Synthetic.
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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ID AATE
                                                                                      RESULT
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Length 18;

Score 11.6; DB 1; Pred. No. 4.8e+02;

0.9%;

Best Local Similarity

Query Match

Sequence 18 BP; 3 A; 3 C; 9 G; 3 T; 0 other;

Example, Page 7; 26pp; English.

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for the treatment of inflammatory bowel disease (IBD) or related disease that treatment of inflammatory bowel disease (IBD) or related disorders that comprises administering a compound that up-regulates the function of IL-9 or the IL-9 receptor. A method for monitoring humans or fragments), by evaluating IL-9 levels in samples taken at different times, and a method for screening for cells expressing the IL-9 receptor. Or the IL-9 receptor and a method for screening for cells expressing the IL-9 receptor to by detecting binding of a specific ligand are also provided. Compounds up-regulating the function of IL-9 receptor can be used the rapeutically (in pharmaceutical compositions, optionally with acceptable carriers) to treat IBD and other related inflammatory disorders. IBDs (which include Crohn's diseases and chronic non-specific inflammatory response to environmental stimuli. Immune responses to antidems are dominated by a Thl or TL2 responses, and evidence suggests that IBDs are dominated by a Thl mediated, antigen induced, inflammatory response. Other related Thl mediated diseases include multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequences shown in AAV41331 to AAV41340 represent exon specific primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       lerosis, diabetes, arthritis, lupus and autoimmune diseases. The me based on the observation that the Th2 response is up-regulated by
                                                                             Interleukin; IL-9; AAFI; asthma associated factor; human; IBD; inflammatory bowel disease; Th2 mediated immune response; lupus; Crohn's disease; chronic non-specific ulcerative colitis; diabetes; multiple sclerosis; arthritis; autoimmune disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating inflammatory bowel diseases, e.g. Crohn's disease - and chronic non-specific ulcerative colitis by administering compounds up-regulating function of interleukin-9 or its receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%; Score 11.6; DB 1; Length 18; ilarity 77.8%; Pred. No. 4.8e+02; Conservative 0; Mismatches 4; Indels
                                        Interleukin-9 (IL-9) gene exon 1 specific lower primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 2 A; 11 C; 2 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 23; 61pp; English.
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                                                                                                                                                                                                                                                                                                                                                                     97US-0994986.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   (MAGA-) MAGAININ PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nicolaides NC;
06-OCT-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-377404/32.
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Les 14; Conserv
                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                              WO9827997-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Levitt RC,
                                                                                                                                                                                                                                                                                     02-JUL-1998
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                                                                                                                                                                                  Synthetic
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the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VBGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatorial arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes nucleic acid molecules which modulate
                                            Vascular endothelial growth factor receptor; VBGF receptor; flt-1; flt-1; ADM; hammerhead ribozyme; halirpin ribozyme; oleavage; tleavage; tleavage; trumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 18;
               Mouse flt-1 VEGF receptor hairpin ribozyme substrate #87.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                                                                                           Stinchcomb D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.9%; Score 11.6; DB 1; 77.8%; Pred. No. 4.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                           Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 188; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      210 CGAAAGCCAGCGAGCTCC 227
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95US-0005974
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(RIBO-) RIBOZYME PHARM INC.
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                                                                                                           foetal liver kinase 1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14; Conservative
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Triticum aestivum.
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABBB399), a sequence having 65% sequence identity to (SI), (SI), acids (SI), applicably conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) concoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples
                                                                                                                                                                                                                                                                                                                                                                                                                                           of the invention.
Note: The present sequence did not form part of the printed
specification, but is based on sequence information supplied to Derwent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, POSHL 1, SH3 domain, POSH-like signalling protein 1; oncogene;
Rho GTPase; signal transduction; gene expression; cancer; vaccine;
                                                                                    Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 11.4; DB 1; Length 17; 92.3%; Pred. No. 4.9e+02; Live 0; Mismatches 1; Indels
                                                                                                                                                     Example 2; SEQ ID NO 220; 60pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human POSHL1 scanning oligonucleotide SEQ ID NO 221.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 2 A; 7 C; 4 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         by the European Patent Office.
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2001WO-US00667.
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2001WO-US00669.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12; Conservative
                                                           WPI; 2002-684061/74.
(AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
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30-JAN-2001; 2
30-JAN-2001; 2
30-JAN-2001; 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABV89508/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   염
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ò
                                                                                                                                                                                                                                                              The present sequence represents a primer for the wheat viviparous 1 (VP1) gene, which keeps mature seeds dormant and inhibits germination. The present invention describes genes which are homologues of the maize Viviparous 1 gene, obtained from oat Avena fatua and wheat which encode polypeptides designated afVP1 and taVP1 respectively. The VP1 activity maintain or impose sufficient intensity and duration and can be used to avoid pre-harvest sprouting (PHS) before harvest. The products can be used in the production of transformed crop plants having desirable primary or secondary dormancy, or after-ripening properties, and in particular may be resistant to PHS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human, POSHL 1, SH3 domain, POSH-like signalling protein 1, oncogene, Rho GTPase; signal transduction, gene expression, cancer, vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.9%; Score 11.6; DB 1; Length 19; 77.8%; Pred. No. 5e+02; ive 0; Mismatches 4; Indels
                                                                                                                                                                                        New isolated oat and wheat VP1 genes, used, e.g. to impose sufficient dormancy to avoid pre-harvest sprouting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human POSHL1 scanning oligonucleotide SEQ ID NO 220.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 3 A; 4 C; 11 G; 1 T; 0 other;
                                                                                                                            Gale MD, Holdsworth MJ;
                                                                                                                                                                                                                                       Claim 56; Page 89; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          721 CAGCAGCAGGGGCCTGG 738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ceccaecaecareceaec 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABV89507 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001WO-US00666.
2001WO-US00667.
2001WO-US00668.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2001WO-US00669.
2001WO-US00670.
2001US-0864761.
2001US-0328205.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene therapy; transgenic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001WO-US00663.
                                                                                              BIOSCIENCE LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-JAN-2002; 2002EP-0001165.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2001WO-US00665.
                                                               97GB-0020060.
                               98WO-GB02835
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 77.8 tes 14; Conservative
                                                                                                                                                        WPI; 1999-244424/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                              (PLAN-) PLANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EP1239051-A2
                                                                                                                            Flintham JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JAN-2001;
                                                                19-SEP-1997;
                                 18-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 685
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABV89507,
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Gaps

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Shannon M;

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The sequences given in AAT80211-382 represent synthetic oligonucleotides which are complementary to a portion of the 5' untranslated region (UTR) of hepatitis C virus (HCV). These sequences may be used in a pharmaceutical composition for the control or prevention of HCV infection. They may be used to inhibit replication or expression of HCV or for detecting the presence of HCV in a sample. They may be used to inhibit HCV replication in a cell and are therefor useful in the treatment of HCV infections such as chronic and acute hepatitis and hepatocarcinoma. This oligo was used in a luciferase assay to determine whether it binds sucessfully to its target.
                                                                                                                                                                                                                                                                                                                              Oligo:nucleotide(s) complementary to HCV 5' untranslated region - used in the treatment and detection of HCV infection, esp. hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepatitis C virus; HCV; hepatocyte infection; non-A hepatitis; non-B hepatitis; acute hepatitis; chronic hepatitis; hepatocellular carcinoms; virucide; cytostatic; antisense therapy; gene therapy; ss; DNA-RNA hybrid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Inhibitory oligonucleotide specific for hepatitis C virus #50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 11.4; DB 1; Length 18; 84.6%; Pred. No. 5.1e+02; tive 1; Mismatches 1; Indels
                                                                                                                                                                                                                        Kilkuskie RE;
Wolfe JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 2 A; 3 C; 10 G; 1 T; 2 U; 0 other;
                                                                                                                                                                                                                        Hamlin HA,
Walther DM,
                                                                                                                                                         (HOFF ) HOFFMANN LA ROCHE & CO AG F. (HYBR-) HYBRIDON INC.
                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 19; Page 31; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                95US-0471968.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96US-021104P.
95US-0471968.
                                                                     96WO-EP02427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97US-0887505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     726 GCAGGGGCCTGG 738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABS65844 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-NOV-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4 GCAGGGGUCCUGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8
Best Local Similarity 84.6
Matches 11; Conservative
                                                                                                                                                                                                                        Goodchild J,
Roberts PC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KILKUSKIE R L.
FRANK B L.
GOODCHILD J.
                                                                                                                                                                                                                                                                                       WPI; 1997-043122/04.
                                                                                                                                                                                                                                                                                                                                                                           and hepato-carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US2002081577-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-JUL-1997;
                                                                     04-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-JUL-1996;
                                                                                                              06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-JUN-1995;
                                                                                                                                                                                                                      Frank BL, G
Roberts NA,
                           12-DEC-1996,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABS65844;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (KILK/) H
(FRAN/) H
(GOOD/) C
(WOLF/) W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABS65844
         8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SL, ABBB399), a sequence having 65% sequence identity to (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to Derwent by the European Patent Office.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                               Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Complementary; 5' untranslated region; UTR; hepatitis C virus; HCV inhibition; replication; expression; detection; chronic hepatitis; acute hepatitis; hepatocarcinoma; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 11.4; DB 1; Length 17; 92.3%; Pred. No. 4.9e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                     Example 2; SEQ ID NO 221; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1..6
/*tag= b
/note= "Phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligo HCV91, targetted to HCV region -1 to -6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 1 A; 7 C; 5 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           a
"2'OMe modified"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT80260 standard; DNA; 18 BP
30-JAN-2001; 2001WO-US00670.
23-MAY-2001; 2001US-0864761.
10-OCT-2001; 2001US-0328205.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        526 CCGGAGGAGCAGC 538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 92.3
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14 CTGGAGGAGCAGC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag=
/note=
                                                                                   (AEOM-) AEOMICA INC
                                                                                                                                                                         WPI; 2002-684061/74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9639500-A2
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Synthetic

AAT80260;

RESULT 687 AAT80260

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Gaps

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The present invention relates to human oligonucleotides comprising a single nucleotide polymorphic site (SNP: AAH88797-AAH89219). The present sequence is one such oligonucleotide. The oligonucleotides can be used in forensics, paternity testing, correlation of polymorphisms with phenotypic traits, genetic mapping of phenotypic traits and marker assisted breeding of animals and crop plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; anglogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; single nucleotide polymorphism; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention describes a polynucleotide (I) of the endothelin (EDNR) fendothelin converting enzyme (ECR)/receptors of EDN and BCE (EDNR) signaling system which is associated with a cardiovascular disease. (I), the gene encoding EDN, BCE or EDNR (II) or a vector (III) expressing (I) or (II) is useful for producing cells capable of expressing a molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful for
                                                                                  New polymorphic sites derived from the human genome are useful to determine sites correlating with phenotypic traits, particularly disease, and also in forensics and paternity testing -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Endothelin converting enzyme 1 (ECE-1) SNP detection primer #63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel polynuclectide of the endothelin/endothelin converting
                                                                                                                                                                                                                                                                                                                                                                                                             Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           Score 11.4; DB 1;
Pred. No. 5.6e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 3 A; 7 C; 6 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
    Thomas D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 6; Page 63; 190pp; English.
                                                                                                                                                                         Claim 95; Page 16; 43pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                             0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-AUG-2001; 2001WO-EP10087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-SEP-2000; 2000EP-0120123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABK94275 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             226 CCTCAGCCTCAGG 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Brinkmann U, Hoffmeyer S;
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8
Best Local Similarity 92.3
Matches 12; Conservative
    Patil N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCTGAGCCTCAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-435060/46.
                                          WPI; 2001-335945/35.
  Chen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200224747-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
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    Au K,
    d
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                     The invention describes synthetic oligonucleotides complementary to a portion of the 5' untranslated region of hepatitis C virus. The oligonucleotides may be used in methods for controlling, preventing, and treating hepatitis C virus infection, in antisense technology and gene therapy, and of detecting the presence of hepatitis C virus in a sample. Hepatitis C virus (HCV) is an enveloped, positive sense, single-stranded RNA virus which infects hepatocytes. HCV is the major cause of non-A, non-B, acute and chronic hepatitis, and has been associated with hepatocellular carcinoma. The invention describes methods and kits for inhibiting replication of HCV, inhibiting the expression of HCV nucleic acid and protein, and for treating HCV infections. This sequence represents a synthetic DNA-RNA hybrid oligonucleotide used for inhibiting HCV replication and expression of HCV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; single nucleotide polymorphic; SNP; forensic science;
paternity testing; phenotypic trait; genetic mapping; animal breeding;
plant breeding; ds.
                                                                                                                                                                                                             Synthetic oligonuclectides complementary to a portion of the 5' untranslated region of hepatitis C virus (HCV), useful for diagnosing and treating HCV infections and hepatocellular carcinoma -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .;
0
                                                                                                        Roberts PC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 11.4; DB 1; Length 18; 84.6%; Pred. No. 5.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 2 A; 3 C; 10 G; 1 T; 2 U; 0 other;
                                                                                                      Wolfe JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
                                                                                                    Goodchild J,
Walther DM;
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replace(10,c)
/*tag= a
                                                                                                                                                                                                                                                                                               Claim 22; Page 11; 74pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     726 GCAGGGGCCTGG 738
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                                                                                  Kilkuskie RL, Fram.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AFFY-) AFFYMETRIX INC.
                 HAMLIN H A.
ROBERTS N A.
WALTHER D M.
  ROBERTS P C.
                                                                                                                                                                       WPI; 2002-537132/57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH89204;
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ROBE/)
                    (HAML/)
                                                              (WALT/)
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Gaps

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variant polypeptide which is associated with a cardiovascular disease.

(II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing

a molecular variant gene computating (I) is useful for identifying and

obtaining a pro-drug or drug capable of modulating the activity of a

molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system

or its gene product, or for identifying and obtaining an inhibitor of

the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE

signaling system or its gene product. The isolated proteins and

colymclectides encoding them are useful for preparation of a

pharmaceutical composition for treating a cardiovascular disease such as

coronary heart disease, hypertension, atherosclerosis, or related to

abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial

CC abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial

CC abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial

CC abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial

CC abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial

CC abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial

CC abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial

CC compusising polynuclectides, gene or a polynucleotide such according to compusising polynucleotides, genes, vectors, polypeptides, antibodies or

CC compusising polynucleotides, genes, vectors, polypeptides, antibodies or

CC compusising polynucleotides, genes, extors, polypeptides, antibodies or

CC compusising polynucleotides, genes, genes, expresents a PCR primer used

CC cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway. Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclearosis; andiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; single nucleotide polymorphism; PCR; primer; ss. Gaps Endothelin converting enzyme 1 (ECE-1) SNP detection primer #64. ö 0.8%; Score 11.4; DB 1; Length 21; 85.7%; Pred. No. 5.8e+02; iive 0; Mismatches 2; Indels Sequence 21 BP; 5 A; 3 C; 11 G; 1 T; 1 other; (EPID-) EPIDAUROS BIOTECHNOLOGIE AG. BP. 19-SEP-2000; 2000EP-0120123. 31-AUG-2001; 2001WO-EP10087 802 CGCTCCCTGCAGCC 815 ABK94276 standard; DNA; 21 (first entry) Brinkmann U, Hoffmeyer S; 18 crcrcccnacaccc s Local Similarity 85.7 1es 12, Conservative WPI; 2002-435060/46. WO200224747-A2 27-AUG-2002 28-MAR-2002 Synthetic. ABK94276; Query Match Best Loca Matches RESULT 691 CCX EX X L X X B X B X QQ ò

Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful

Example 6; Page 63; 190pp; English.

the disease

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signaling system which is associated with a cardiovascular disease. (I), the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing (I) or variant polypeptide which is associated with a cardiovascular disease.

Cr variant polypeptide which is associated with a cardiovascular disease.

(II), (III), the EDN ECE or EDNR polypeptide, or a cell expressing or a molecular variant gene comprising (I) is useful for identifying and obtaining a pro-drug or drug capable of modulating the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product. The isolated proteins and control of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product. The isolated proteins and pharmaceutical composition for treating a cardiovascular disease such as coronary heart disease, hypertension, atherosclerosis, or related to abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial hypercholesterolaemia. The gene or a polymuclectide fragment of the EDN/EDNR signaling system are useful as forensic markers, for creating a transgenic animal and in creation of a solid support creating a transgenic animal and in creation of a solid support comprising polymuclectides, antibodies or comprising polymuclectides, antibodies or comprising polymuclectides, antibodies or comprising polymuclectides, antibodies or comprising polymuclectides, and pharence represents a PCR primer used
                                                                                                                                                                                                                                                                                                                                                                                                    to identify single nucleotide polymorphisms in DNA encoding cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, variant thrombospondin 1, variant thrombospondin 4, SNP; polymorphism, vascular disease, coronary artery disease, forensics; myocardial infarction, atherosclerosis, stroke, venous thromboembolism;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 1 A; 11 C; 3 G; 5 T; 1 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (WHED ) WHITEHEAD INST BIOMEDICAL RES. (MILL-) MILLENNIUM PHARM INC.
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26-JUL-2000; 2000US-0220947.
16-AUG-2000; 2000US-0225724.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                802 CGCTCCCTGCAGCC 815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 CTCTCCCNGCAGCC 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-226749/23
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Best Local Similarity
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Variation
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            8899999999999999999988888
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Page 338

The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, mycoardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification. Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis. Examples; Page 116; 242pp; English. 

Sequence 21 BP; 5 A; 8 C; 7 G; 1 T; 0 other;

0.8%; Score 11.4; DB 1; Length 21; 71.4%; Pred. No. 5.8e+02; tive 0; Mismatches 6; Indels 892 CTGCGGTACAGCGTGGCCCTG 912 21 crecerecaderrescerie 1 Query Match 0.8 Best Local Similarity 71.4 Matches 15; Conservative g ò

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Gaps ;

> RESULT 693 ACA06326,

ACA06326 standard, RNA; 17 BP

ACA06326; 

03-JUN-2003 (first entry)

NFKB sub-unit modulating inozyme substrate #145.

Enzymatic nucleic acid, nuclear factor kappa B; NFKB; inozyme; de-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; cospangeal cancer; stomach cancer; baldder cancer; pancreatic cancer; cervical cancer; stomach cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; doctaxel; cisplatin; methorrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn; s disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection;

Homo sapiens.

US2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-0864785.

92US-0987132. 94US-0245466. 96US-0777916. 94US-0291932 15-AUG-1994; 07-DEC-1992; 18-MAY-1994; 23-DEC-1996;

STINCHCOMB D T. MCSWIGGEN J. DRAPER K G. (/NILS DRAP/) MCSW/

Stinchcomb DT, Mcswiggen J, Draper KG;

WPI; 2003-340953/32

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases

Claim 3; Page 29; 72pp; English.

The intention describes an enzymatic muclear acta molecule (1) wince the regulates expression of a sequence encoding a subunit of muclear factor kappa B (NFKB), where (1) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (1) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG2+ The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodias, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, gemittabine or radiation therapy. The enzymatic and antisense nucleic cond molecules are also useful for treating inflammatory disease such as necessity, autoimmune disease, lupus, multiple solerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury central nervous system (CNS) and myocardial), glomerulosaphitis, especial are electroned in the propertion inflammatory bowel disease or section method and propertion inflammatory bowel disease or invention describes an enzymatic nucleic acid molecule (I) which down infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule.

Sequence 17 BP; 0 A; 11 C; 3 G; 3 U; 0 other;

Gaps . 0 0.8%; Score 11.2; DB 1; Length 17; 11.2%; Pred. No. 5.3e+02; ve 0; Mismatches 3; Indels 81.2%; Query Match
Best Local Similarity 81.2'

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466 AGCCTGCAGGGGAGG 481 17 Aggcggcacacacaca

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AAQ26549 standard; DNA; 18 RESULT 694 AAQ26549

BP.

AAQ26549; 

08-JAN-1993

(first entry)

Control probe #4 for caucosoid RING11 gene.

immunosuppressants; immunoenhancers; treatment; diagnosis; screening; immune disorders; transporter peptides; proteasome complex; MHC class I molecules; HLA; antigen processing; antigen presentation; autoimmune disease; ankylosing spondylitus; prenatal diagnosis; polymerase chain reaction; sc

Synthetic.

WO9211289-A

09-JUL-1992.

91WO-GB02278 19-DEC-1991; 90GB-0027520 91GB-0019711 19-DEC-1990; 16-SEP-1991; (IMCR ) IMPERIAL CANCER RES TECHNOLOGY.

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The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 460 GTCAGCAGCCTGCAGGGG 478
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                                                                                                                                                                                                                     AAX33259 standard; DNA; 20 BP.
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97JP-0254250.
97JP-0299407.
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                                                                                          Best Local Similarity 81.8 Matches 9; Conservative
                                                                                                                               731 GGGCCTGGCTG 741
                                                                                                                                                       12
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Ogawa S, Tagashira S;
                                                                                                                                               WPI; 1999-243621/20.
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15-OCT-1997;
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                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                               Query Match
                                                                                                                                                                                                                                             AAX33259
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ABS74296/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
                                                                                                                                                                                                                                             Gaps
                                                                                                                                     This probe was used together with AAQ26546-51 to analyse caucosoid controls by oligonucleotide typing, whilst investigating RING 11 polymorphisms - see AAQ26544,5.
                                                           DNA encoding RING4, RING10, RING11 AND RING12 proteins - for treatment and diagnosis of immune disorders and screening of new immunosuppressants and immuno:enhancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human, ribozyme, short interfering RNA, siRNA, HER2, K-Ras,
enzymatic nucleic acid, H-Ras, N-Ras, HIV; cytostatic, anti-HIV;
anti-rheumatic, cancer, AIDS, ss.
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0
                                                                                                                                                                                                                    Length 18;
                                                                                                                                                                                                                                          3; Indels
                                                                                                                                                                                                                0.8%; Score 11.2; DB 1;
81.2%; Pred. No. 5.5e+02;
              Trowsdale J;
                                                                                                                                                                                        Sequence 18 BP; 3 A; 6 C; 6 G; 3 T; 0 other;
                                                                                                                                                                                                                                         0; Mismatches
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                                                                                                              Example 2; Page 40; 101pp; English.
              Powis SH,
                                                                                                                                                                                                                                                                                                                                                                                                                                Human H-Ras DNAzyme target #357.
                                                                                                                                                                                                                                                                 631 CTCCAGGAGCTCTGCA 646
                                                                                                                                                                                                                                                                                                                                                       ABZ61566 standard; RNA; 17 BP
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06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
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2001US-318471P.
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                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                       13; Conservative
             Kelly AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-140484/13.
                                   WPI; 1992-250030/30
                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200297114-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                        21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-DEC-2002
           Glynne R,
                                                                                                                                                                                                                                                                                                                                                                               ABZ61566;
                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                              RESULT 695
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes DNA which participates in the regulation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of expression of PEBP2 alpha A gene. The DNA produces a regulation protein with the activity of promoting bone formation and can serve as promoter for prevention and treatment of bone diseases including osteoporosis. The present sequence represents a PCR primer used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PEBP2 alpha A gene expression regulating DNA PCR primer SEQ ID NO:16.
                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA regulating expression of PEBP2 alphaA gene to produce regulator protein, useful as promoter for prevention or/and treatment of bone diseases e.g. osteoporosis
                                                                                                                                  ó.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ..
                                                          0.8%; Score 11; DB 1; Length 17;
81.8%; Pred. No. 5.7e+02;
tive 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PEBP2 alpha A gene; expression; regulation; bone disease; osteoporosis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 11, DB 1; Length 20;
Pred. No. 6.4e+02;
0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Katsumata T, Nakatsuka M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 11 C; 4 G; 2 T; 0 other;
Sequence 17 BP; 2 A; 4 C; 8 G; 3 U; 0 other;
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ABS74296 standard; DNA; 20

ABS74296;

(first entry) 09-DEC-2002

Auman calcium channel alpha2delta SSCP PCR primer #20.

Human; ss; primer; calcium channel alpha2delta; splice isoform; CACNA2D2; gene therapy, Lambert-Eaton myssthenic syndrome, LEMS; PCR, autoimmune disease; epilepsy; migraine; episodic ataxia; cancer; stroke; brain trauma; Alzheimer s' disease; multinfarct dementia; convulsion; Korsa-Koff's disease; amyotrophic lateral sclerosis; seizure; Huntington's disease; ammesia; cardiac arrhythmia; sugina pectoris; hypoxia; ischaemia; myocardial infarction; congestive heart failure; muscular dystrophy; hypertension; chromosome 3p21.3; lung cancer; breast cancer; preneoplastic lesion; hyperplasia; dysplasia; carcinoma; SSCP; single strand change polymorphism.

Homo sapiens

US6441156-B1

27-AUG-2002.

99US-0470443 22-DEC-1999; 98US-114359P 30-DEC-1998;

(USSH ) US DEPT HEALTH & HUMAN SERVICES.

Gao Duh F, Minna JD, Sekido Y, Wei M, Latif F, WPI; 2002-730574/79. Lerman MI,

Novel purified nucleic acid sequence encoding human calcium channel alpha2delta subunit protein, useful for detecting, preventing and treating cancer, stroke, brain trauma, Huntington's disease, myocardial infarction

Example 7; Column 46; 77pp; English.

CCROMAZD2 gene which encodes human calcium chamnel alphazdelta-2 subunit protein) comprising a fully defined alphazdelta splice isoform 1, 2 or a nucleic acid sequence, or its complement and the encoded proteins.

Also include are: (1) a method of producing a calcium chamnel protein comprising a recombinant expression vector comprising the calcium chamnel protein into calcium chamnel protein, comprising carrying out the method of (1), but with one chamnel proteins, comprising carrying out the method of (1), but with one or more than one expression vector comprising calcium chamnel protein or comprising carrying patients with lambert-Eaton myasthenic sequences encoding the splice variants. CACNAZDZ nucleic acid is useful for treating patients with lambert-Eaton myasthenic syndrome (LBMS) (an autoimmune disease) and for identifying compounds the splice variants with lambert-Eaton myasthenic syndrome (LBMS) (an autoimmune disease) and for identifying compounds useful for treating other diseases associated with abnormal calcium channel protein activity (e.g. epilepsy, migraine, episodic ataxia, channel protein activity (e.g. epilepsy, migraine, episodic ataxia, channel protein activity (e.g. epilepsy, migraine, episodic ataxia, setures, hypoxic damage to the cardiovascular system, ischaemic damage to the cardiovascular system, ischaemic damage to the cardiovascular system, inchained and probes for detecting presence of nucleic acid is sequence encoding at least a portion of calcium channel protein, in detection, identification and isolation of alpha2delta sequences (e.g. lung cancer and breast cancer) and cancer (e.g. lung cancer and breast cancer) and present is an SSCP (single strand change polymorphism) PCR primer The present is an SSCP (single strand change polymorphism) PCR primer and protein cancer (e.g. hyperplasia, dysplasia, cariar calcium cancer (e.g. hyperp The invention relates to a purified nucleic acid sequence (referred as

ö Gaps used to detect polymorphisms in sequences encoding a human calcium channel alpha2delta splice isoform protein. ö Length 20; 5; Indels Score 11; DB 1; I Pred. No. 6.4e+02; Sequence 20 BP; 4 A; 4 C; 8 G; 4 T; 0 other; 0; Mismatches 618 CTTCAGGGACCAGCTCCAG 636 19 CICCIGIGACCAICACCAG 1 0.8%; 14; Conservative Similarity Query Match Best Local SXSS à 쉽

RESULT 698 ABS74306,

ABS74306 standard; DNA; 20

ABS74306;

(first entry) 09-DEC-2002 Human calcium channel alpha2delta SSCP PCR primer #30.

Human; ss; primer; calcium channel alpha2delta; splice isoform; CACNA2D2; gene therapy; Lambert-Eaton myasthenic syndrome; LEMS; PCR; autoimmune disease; epilepsy; migrains; episodic ataxia; cancer; stroke; brain trauma; Alzheimer: s disease; multiinfarct dementia; convulsion; Korsa-Koff's disease; amyotrophic lateral sclerosis; seizure; Huntington's disease; amyotrophic lateral sclerosis; seizure; Huntington's disease; amnesia; cardiac arrhythmia; angina pectoris; muscular dystrophy; hypertension; chromosome 3p21.3; lung cancer; breast cancer; preneoplastic lesion; hyperplassia; dysplasia; carcinoma; SSCP; single strand change polymorphism.

Homo sapiens

US6441156-B1

27-AUG-2002.

99US-0470443. 22-DEC-1999; 98US-114359P. 30-DEC-1998;

(USSH ) US DEPT HEALTH & HUMAN SERVICES.

Sekido Y, Latif F, Wei M, Duh F, Minna JD, Lerman MI,

WPI; 2002-730574/79.

Novel purified nucleic acid sequence encoding human calcium channel alpha2delta subunit protein, useful for detecting, preventing and treating cancer, stroke, brain trauma, Huntington's disease, myocardial infarction

Example 7; Column 47; 77pp; English.

The invention relates to a purified nucleic acid sequence (referred as CACNA2D2 gene which encodes human calcium channel alpha2delta-2 subunit protein) comprising a fully defined alpha2delta splice isoform 1, 2 or 3 nucleic acid sequence, or its complement and the encoded proteins.

Also include are: (1) a method of producing a calcium channel protein which involves introducing a recombinant expression vector comprising the CACNA2D2 nucleic acids and encoding the calcium channel protein, into a cultured host cell under conditions such that the host cell expresses the amino acid sequences; and (2) a method for co-expressing calcium channel proteins, comprising out the method of (1), but with one or more than one expression vector comprising one or more nucleic acid sequences encoding the splice variants. CACNA2D2 nucleic acid is useful sequences encoding the splice variants. CACNA2D2 nucleic acid is useful for treating patients with lambert-Eaton myasthenic syndrome (LEMS) (an autoimmune disease) and for identifying compounds

channel protein activity (e.g. epilepsy, migraine, episodic ataxia, cancer, stroke, brain trauma. Alzheimer's disease, multinfarct dementia, CC cancer, stroke, brain trauma. Alzheimer's disease, multinfarct dementia, CXOISA-KOff's disease, amyotrophic lateral sclerosis, convulsions, control of disease, ammeeia, cardiac arrhythmia, angina pectoris, hypoxic damage to the cardiovascular system, ischaemic damage to the cardiovascular system, ischaemic damage to the cardiovascular system, congestive heart failure, muscular dystrophy and hyportension) CACNA2D2 nucleic acid is useful as primers and probes for detecting presence of nucleic acid is captance encoding at least a portion of calcium channel protein, in detection, identification and isolation of alpha2delta sequences diagnosing and typing of preneoplasias and cancers, since genetic cid supption of 3p21.3 region (in which the alpha\_2delta gene is located) is common in cancer (e.g. lung cancer and breast cancer) and preneoplastic lesion (e.g. lupperplasia, dysplasia, carcinoma in situ). The present is an SSCP (single strand change polymorphism) PCR primer channel alpha2delta encoding a human calcium channel alpha2delta and colium channel alpha2delta carcinoma in situ). channel alpha2delta splice isoform protein. 

Sequence 20 BP; 4 A; 4 C; 8 G; 4 T; 0 other;

.; 0 0.8%; Score 11; DB 1; Length 20; 73.7%; Pred. No. 6.4e+02; 0; Mismatches Conservative Local Similarity les 14; Conserv Query Match Best Loca Matches à

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RESULT 699 AAI66616

AAI66616 standard; DNA; 20 BP

AA166616;

(first entry) 07-JAN-2002

Rat leukotriene B4 receptor JULF2 DNA amplifying PCR primer.

Leukotriene receptor; leukotriene B4; inflammatory disease; rat; JULF2; bronchitis; dermatitis; psoriasis; ulcerative colitis; rheumatoid arthritis; edema; PCR primer; ss.

Rattus norvegicus,

WO200170815-A1.

27-SEP-2001

15-MAR-2001; 2001WO-JP02060.

2000JP-0078992. 2000JP-0187978. 21-MAR-2000; 22-JUN-2000;

(YAMA ) YAMANOUCHI PHARM CO LTD.

New polypeptide for screening for compounds which treat inflammatory diseases such as bronchitis, dermatitis, psoriasis, ulcerative colitis, rheumatoid arthritis, and edema comprises the leukotriene B4 receptor -WPI; 2001-611487/70.

Ohishi T;

Saito T,

Takasaki J,

Matsumoto M,

Kamohara M,

Example 10; Page 47; 55pp; Japanese.

The invention provides a leukotriene receptor, which binds leukotriene B4 and polynucleotides encoding the leukotriene B4 receptor. The receptor can be expressed by standard recombinant methodology. Pharmaceutical compositions containing materials which modify the receptor activity, other than 4-octyloxybenzene carboximidoamide are used for treating and preventing inflammatory disease. The materials detected by screening the 

ò Gaps receptor (JULF2) are useful for treating diseases such as bronchitis, dermacitis, psoriasis, ulcerative colitis, rheumatoid archritis, and edema. Sequences AA166614-13 represent PCR primers for amplifying a rat leukotriene B4 receptor JULF2 DNA. 0 Score 11; DB 1; Length 20; Pred. No. 6.4e+02; 0; Mismatches 5; Indels Sequence 20 BP; 5 A; 11 C; 4 G; 0 U; 0 other; 585 CCGTCTGCCCCCCACCAGC 603 cadocadaccoccadocado 20 L Similarity 73.7%; 14; Conservative 2 Query Match Local Matches 88888888 ð a

ABK15887 standard; DNA; 20 BP. (first entry) 21-MAY-2002 ABK15887; RESULT 700 ABK15887,

Hairy-enhancer of split-1; HES-1; real-time PCR; primer; ss; multiple sclerosis; rheumatoid arthritis; diabetes; organ transplant; asthma, allergy; autoimmunity; graft rejection; tumour; cytostatic; Notch signal modulator; T-cell mediated disease; infectious disease; human immunodeficiency virus; HIV; virucide; hepatotropic; protozoacide; HES-1 (hairy-enhancer of split-1) forward PCR primer DNA sequence. neuroprotective; cancer.

Unidentified

WO200212890-A2

14-FEB-2002.

03-AUG-2001; 2001WO-GB03503.

04-AUG-2000; 2000GB-0019242.

(LORA-) LORANTIS LTD

Champion BR; Dallman MJ, WPI; 2002-217232/27. Lamb JR, Hoyne GF, 

T-cell Monitoring the immune system for prevention and/or treatment of mediated diseases e.g. allergy, autoimmunity or cancer, involves detecting modulation of Notch signalling

Disclosure, Fig 13; 75pp; English.

The present invention relates to a new method for monitoring the immune system that involves detecting modulation of Notch signalling. The method of the invention can be used for monitoring the immune system such as detecting or monitoring T-cell activation or inactivation, immunological colerance or activity, monitoring the efficacy of immunotherapy and for detecting or monitoring the reactivity of a T-cell to an antigen e.g. for detecting increased or decreased reactivity of a T-cell to an antigen and detecting tolerisation of a T-cell to an antigen and detecting tolerisation of a T-cell to an antigen. The method is used in the complex the antigen is self or foreign antigen. The method is used in the prevention and/or treatment of T-cell mediated diseases such as asthma, or prevention and/or treatment of T-cell mediated diseases such as asthma, alleray, autoimmunity, graft rejection, tumour induced aberrations to the prevention and/or treatment of T-cell mediated diseases caused by e.g. Cytomegalovirus, beauchomonas, Toxoplasma, Microfilariae, Helminths, Mycobacteria, human immunodeficiency virus (HIV), plasmodium species, Echinococcus, communodeficiency virus (HIV), plasmodium species, Echinococcus, comethod is also used for the transplantation. The present assay cuthritis, diabetes and for organ transplantation. The present assay method provides a much more objective measure of the effectiveness of

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Gaps

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therapy than the rather subjective symptoms-based measures which are often used at present. The ability to detect an immune response could be used in identifying the cause of an allergic reaction by monitoring the activity of the immune system in the presence of different potential allergens. The assay could be used to check for successful immunisation against a given disease antigen. The present nucleic acid sequence represents the human HES-1 (hairy-enhancer of split-1) forward PCR primer that was used in the invention with the HES-1 reverse PCR primer (ABK15888) and the HES-1 probe (ABK09925) for real-time PCR of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This 3' primer and a 5' primer (see AAV47651) are based on an area of non-homology between murine protein tyrosine kinase 2 (PYK2) and focal adhesion kinase (FAK) that is adjacent to the C-terminus of the kinase domain. They were used in a PCR amplification of CDNAs of mouse osteoblastic MB1.8. The PCR product (700 bp) was used as a FAK-specific probe to isolate mouse FAK CDNA. The invention relates to new nucleic acid (see AAV47653) encoding mouse PYKZ (see AAW61196), a member of the FAK family. PYKZ can be used in a claimed method for identifying specific modulators of PYKZ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid encoding murine protein tyrosine kinase 2 and cells expressing the recombinant kinase - used to identify specific modulators, potentially useful for controlling the level of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protein tyrosine kinase 2; PYK2; mouse; cell adhesion kinase-beta; related adhesion focal tyrosine kinase; focal adhesion kinase; platelet; PCR; primer; ss.
                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                   Score 11; DB 1; Length 20; Pred. No. 6.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse focal adhesion kinase cDNA 3' PCR primer.
                                                                                                                                                                                                 Sequence 20 BP; 5 A; 5 C; 6 G; 4 T; 0 other;
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                                                                                                                                                                                                                                                                                                         435 GTTCAGAAAGTTGCTGAAG 453
                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                                                            GTTCATGCACTCGCTGAAG 2
                                                                                                                                                                                                                                  Query Match 0.8%;
Best Local Similarity 73.7%;
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                AAV47652 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-447214/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-DEC-1998
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                                                                                                                                                               HES-1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                                                                                                                                                                                                                                                                                                                                          20
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                                                                                                                                                                                                                                                                                                                                                                                            RESULT 701
AAV47652/c
ID AAV476
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This oligonucleotide is based on a non-homologous region, found adjacent to the C-terminal of the kinase domain, of murine protein tyrosine kinase 2 (PYK2) and focal adhesion kinase (FAK) sequences. It was used as a 3 primer, together with a 5' primer (see AAV45666), in the PCR amplification of mouse osteoblastic MB1.8 cell cDNA. The PCR product was used as a FAK-specific probe to isolate full-length FAK cDNA. FAK shows homology to murine PYK2 (see AAW6568), another cell adhesion-dependent kinase. Agents that bind to and modulate PYK2 are isolated using methods of the invention, and are useful in treating osteoporosis and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying agents that bind and modulate protein tyrosine kinase 2 - useful for inhibiting migration, adhesion or activity of monocytic cells, particularly for treatment and prevention of osteoporosis and inflammation
                                Gaps
                                                                                                                                                                                                                                                                               Focal adhesion kinase; protein tyrosine kinase 2; PYK2 gene; mouse; podosome; related adhesion focal tyrosine kinase; cell adhesion kinase; ligand; moncyte; osteoporosis; inflammation; therapy; PCR; primer; ss.
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0.8%; Score 11; DB 1; Length 21; 73.7%; Pred. No. 6.6e+02; ve 0; Mismatches 5; Indel8
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                                                                                                                                                                                                                                                   Focal adhesion kinase 3' PCR primer.
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                                                           1019 GATGGTGCCAAAGTGCAGC 1037
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                                                                                                                                                               21
                                                                                                                                                                                                                       (first entry)
Query Match 0.8
Best Local Similarity 73.7
Matches 14; Conservative
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                                                                                                                                                              AAV49607 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rodan GA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-447250/38.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                           AAV49607;
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                                                                                                                                  RESULT 702
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RESULT 703 ABL31720/c

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Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when
                                                                                                                                                                                                                                    Nishida M;
                                                                           Human; human leukocyte antigen; HLA; genotype; polymozphism;
immunogenetic; transplantation; genetic disease; ss.
                                                                                                                                                                                                                                    Moriya S,
                                                        Human HLA genotyping oligonucleotide SEQ ID NO 1209,
                                                                                                                                                                                                                                 Ichihara T, Matsumura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence 17 BP; 5 A; 4 C; 6 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                             Claim 10; Page 322; 345pp; Japanese.
                                                                                                                                                                01-JUN-2001; 2001WO-JP04662.
                                                                                                                                                                                   01-JUN-2000; 2000JP-0164798
ABL31720 standard; DNA; 17
                                                                                                                                                                                                                                                                                           transplanting between them
                                      (first entry)
                                                                                                                                                                                                     (NISN ) NISSHINBO IND INC
                                                                                                                                                                                                                (SYST-) SYSTEM RES INC
                                                                                                                                                                                                                                                     WPI; 2002-122074/16.
                                                                                                                                                                                                                                  Kagiya T,
                                                                                                                          WO200192572-A1.
                                                                                                        Homo sapiens
                                     21-MAR-2002
                                                                                                                                            06-DEC-2001
                   ABL31720;
                                                                                                                                                                                                                                  Inoko H,
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The invention relates to a typing kit for judging human leukocyte antigen (HIA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleorides (ABL30512-ABL31809) originating in the sequences of genes e.g. belonging to HIA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved mucleic acids relating to gene polymorphisms. The method is useful for judging HIA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of pancreas, largerhans islet in pancreas and cornea, susceptibility diancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals.

Query Match

0.8%; Score 10.8; DB 1; Length 17; 95.7%; Pred. No. 6.1e+02; ve 0; Mismatches 2; Indels 85.7%; 974 TCACTIGACCAGIC 987 Conservative Local Similarity les 12; Conserv Best Loca Matches

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rcacreseccaere 4 17

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RESULT 704 ACA07770 ACA07770;

ACA07770 standard; RNA; 17 BP

NFKB sub-unit modulating zinzyme substrate #169.

(first entry)

03-JUN-2003

Enzymatic nucleic acid, nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; cesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; 

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lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphamide; doxrubin; fluorouracil carboplatin; edatrexate; gemcitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection;
```

Homo sapiens.

US2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-0864785.

.5-AUG-1994;

94US-0291932. 92US-0987132. 94US-0245466. 96US-0777916. 07-DEC-1992; L8-MAY-1994;

23-DEC-1996; 

(STIN/) STINCHCOMB D T. (MCSW/) MCSWIGGEN J.

DRAPER K G.

(DRAP/)

Draper KG; Stinchcomb DT, Mcswiggen J,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases

Claim 3; Page 40; 72pp; English.

regulates expression of a sequence encoding a subunit of nuclear factor.

Kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating FEL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, centiables are also useful for treating inflammatory disease such as charactering and antisense nucleic acid molecules are also useful for treating inflammatory disease such as charactering archivity and antisease such as charactering activity and antisease such as charactering and antisense such as charactering activity and antisease such as charactering and antisense such as charactering activity and antisease such as charactering and antisense such as charactering activity and antisease such as charactering activity and antisease such as charactering activity and antisease such as charactering and antisense such as charact invention describes an enzymatic nucleic acid molecule (I) which down rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (Central nervous system (CNS) and mycoardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule.

Sequence 17 BP; 2 A; 6 C; 6 G; 3 U; 0 other;

Gaps .. Length 17; Indels 0.8%; Score 10.8; DB 1; 14.3%; Pred. No. 6.1e+02; ve 3; Mismatches 2; 64.38; 9; Conservative Best Local Similarity Query Match Matches

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820 GTCCTGATGCAGCT 833

à dd

4 GCCCUGCUGCAGCU 17

us09904568-3.rng

hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesteroleamia; forensic marker; transgenic animal; solid support; cardicvascular regulator; SNP; single nucleotide polymorphism; PCR; primer; ss.

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Drosophila circadian rhythm period gene; circadian clock gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human genes. The RIGHI polypeptides act as regulators of circadian rhythms. The identification of RIGHI as a putative circadian clock gene provides a useful tool to explore the molecular mechanism of the mammanian circadian machinery. Using interaction screening approaches, it should be possible to find interaction screening perhaps in the form of a Drosophila Timeless orthology. Furthermore, promoter analyses of the RIGHI genes should uncover how light cues and possibly other environmental stimuli, regulate the expression of this gene. Targeted disruption of the might gene using stem cell technology, may provide a valuable model system to study the various physiclogical and pathological aspects of disrupting circadian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primers AAX26903-04 were used for STS-PCR mapping of RIGUI nucleic acids. RIGUI is a gene corresponding to the Drosophila circadian rhythm period gene. The specification describes both mouse and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Indels
                                                                                                              Primer used for STS-PCR mapping of RIGUI nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 10.8; DB 1;
85.7%; Pred. No. 6.8e+02;
ative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New isolated mammalian circadian rhythm genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 7 C; 5 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                              Sun ZS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 30; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                             Lee C,
                  AAX26904 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                98WO-US18755
                                                                                                                                                                                                                                                                                                                             97US-0065957
                                                                                                                                                                                                                                                                                                                                              97US-0058256
                                                                                (first entry)
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Best Local Similarity 85.7
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                           Eichele G,
                                                                                                                                                                                                                                                                                                                                                                           (RERE-) RES DEV FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-229221/19.
                                                                                                                                                                                                                                                                                                                                              09-SEP-1997;
                                                                                                                                                                                                                               W09912952-A1
                                                                                                                                                                                                                                                                                              09-SEP-1998;
                                                                                                                                                                                                                                                                                                                             04-NOV-1997;
                                                                                23-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                           Albrecht U,
                                                                                                                                                                                                                                                              18-MAR-1999
                                                                                                                                                              Drosophila
                                                                                                                                                                                              Synthetic.
                                                 AAX26904;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chythms
                                                                                                                                               RIGUI;
AAX26904/
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Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful for

Example 6; Page 63; 190pp; English.

treating the disease

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Hoffmeyer S;

Brinkmann U,

WPI; 2002-435060/46.

31-AUG-2001; 2001WO-EP10087.

WO200224747-A2

Synthetic.

28-MAR-2002.

19-SEP-2000; 2000EP-0120123

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The invention describes a polynucleotide (I) of the endothelin (EDNR) /endothelin converting enzyme (ECB)/receptors of EDN and ECE (EDNR) (C (EDN)/endothelin converting enzyme (ECB)/receptors of EDN and ECE (EDNR) (C (EDN)/endothelin spassociated with a cardiovascular disease. (I), the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing a molecular variant polypeptide which is associated with a cardiovascular disease. (II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing and a molecular variant gene comprising (I) is useful for identifying and cotaling a product, or for identifying and obtaining an inhibitor of the activity of a molecular variant of a polypeptide of the activity of a molecular variant of a polypeptide of the activity of a collectular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product. The isolated proteins and correctlides encoding them are useful for preparation of a polymential angiogenesis or fatty acid metabolism e.g. diabetes and familial hypercolaster comparison, atherosclerosis, or related to abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial hypercolaster comparison of a solvated proteins and correcting a transgenic animal and in creation of a solid support comparising polymucleotides, genes or expresents a PCR primer used to comprising polymucleotides, genes vectors, polypeptides, and tibodies or host cells of the invention. This sequence represents a PCR primer used to identify single nucleotide polymorphisms in DNA encoding cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Endothelin converting enzyme 1 (ECE-1) SNP detection primer #62.
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 10.8; DB 1;
15.7%; Pred. No. 7e+02;
.ve 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 5 A; 3 C; 12 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Query Match
Best Local Similarity 85.,,,
Best Local 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK94274 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CGCTCCCTGCAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTCTCCCCGCAGCC
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Gaps

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Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease;

Endothelin converting enzyme 1 (ECE-1) SNP detection primer #61.

(first entry)

27-AUG-2002

MAXEXEX MAXEX MAXE

ABK94273;

ABK94273 standard; DNA; 21 BP.

RESULT 706

ABK94273

333 TCCTGGTGATAGTC 346

δ

g

15 rccredacarecre 2

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Human KTOMla portion (ABQ63232) probe # 348.

(first entry)

20-AUG-2002

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Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; BDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; andiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; single nucleotide polymorphism; PCR; primer; ss.
                                                                                                                                                                                                                enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful for treating the disease.
                                                                                                                                                                                                      Novel polynucleotide of the endothelin/endothelin converting
                                                                                                                                                   (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                                                Example 6; Page 63; 190pp; English.
                                                                                                                   31-AUG-2001; 2001WO-EP10087.
                                                                                                                                   19-SEP-2000; 2000EP-0120123
                                                                                                                                                                     Brinkmann U, Hoffmeyer S;
                                                                                                                                                                                      WPI; 2002-435060/46.
                                                                                  WO200224747-A2.
                                                                                                   28-MAR-2002
                                                                 Synthetic.
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cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway. creating a transgenic animal and in creation of a solid support comprising polynucleotides, genes, vectors, polypeptides, antibodies or host cells of the invention. This sequence represents a PCR primer used identify single nucleotide polymorphisms in DNA encoding Sequence 21 BP; 1 A; 12 C; 3 G; 5 T; 0 other;

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Gaps
                                       ..
0
 0.8%; Score 10.8; DB 1; Length 21;
85.7%; Pred. No. 7e+02;
tive 0; Mismatches 2; Indels
                 Local Similarity 85.7
es 12; Conservative
Query Match
                 Best Loca
Matches
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802 CGCTCCCTGCAGCC 815 CTCTCCCCCCAGCC 17

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ABQ63635 standard; DNA; 17 BP. ABQ63635; ABQ63635

RESULT 708

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The invention relates to a novel isolated nucleic acid encoding human invention has cytostatic activity. The nucleotide may have a use in gene therapy. The KTOM1 (Midney tumour oversxpressed membrane) protein. The protein of the therapy. The KTOM1 nucleic acids may be used to diagnose, treat or monitor a disease caused by altered expression of human KTOM1.

Compositions comprising the nucleic acids, proteins or antibodise may be used to treat subjects having defects in KTOM1 which can manifest as cancer of the kidney, as well as a disorder of liver, bone marrow, brain, heart, lung, kidney, colon, skeletal muscle, testis, uterus and placenta function. The sequence represents a probe used in the invention to scan the nt 1-1001 portion of human KTOM1a (ABQ631232).
                                                        Human, KTOMla; KTOMl; kidney tumour overexpressed membrane; cytostatic; gene therapy; cancer; kidney; liver; bone marrow; brain; heart; lung; kidney; colon; skeletal muscle; testis; uterus; placenta; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acids encoding the protein, useful for treating subjects having defects in KTOM1 which can manifest as cancer of the kidney, or as a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       overexpressed membrane (KTOM1) protein and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 10.6; DB 1; Length 17; 76.5%; Pred. No. 6.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 203; 418pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disorder of e.g., liver or bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1316
                                                                                                                                                                                                                                                                                                                30-JAN-2001; 2001W0-US00663.
30-JAN-2001; 2001W0-US00663.
30-JAN-2001; 2001W0-US00664.
30-JAN-2001; 2001W0-US00666.
30-JAN-2001; 2001W0-US00666.
30-JAN-2001; 2001W0-US00666.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
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                                                                                                                                                                                                                     21-SEP-2001; 2001WO-US29656
                                                                                                                                                                                                                                                                                  2000GB-0024263
2001WO-US00661
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New human kidney tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-479509/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                      WO200224750-A2.
                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                          30-JAN-2001;
                                                                                                                                                                                                                                                    21-SEP-2000;
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                                                                                                                                                                                                                                                                                    04-OCT-2000;
                                                                                                                                                                                       28-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zhang J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 709
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셤
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Human sentrin phosphorothioate antisense oligonucleotide SEQ ID NO:47.

(first entry)

03-FEB-2000

AAZ35905;

AAZ35905 standard; DNA; 18 BP.

AAZ35905

Human, sentrin, antisense oligonucleotide, phosphorothioate, inhibition, modulation, expression, diagnosis, ss.

1..18 /\*tag= a /note= "phosphorothioate linkages"

98US-0213768

17-DEC-1998;

16-NOV-1999.

US5985664-A.

17-DEC-1998;

(ISIS-) ISIS PHARM INC.

Cowsert LM;

Baker BF,

WPI; 2000-022284/02.

Location/Qualifiers

Key modified\_base

Homo sapiens

Synthetic

```
Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;
                                                                                                                                                                                                                                                                                                                                   Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 10.6; DB 1; Length 17; 76.5%; Pred. No. 6.5e+02; Ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                         Example 2; SEQ ID NO 494; 60pp + Sequence Listing; English.
                           Human POSHL1 scanning oligonuclectide SEQ ID NO 494.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 3 A; 4 C; 9 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the European Patent Office.
                                                                   Jene therapy; transgenic; ss.
                                                                                                                                                                                                                 2001WO-US00668.
2001WO-US00669.
2001WO-US00670.
                                                                                                                                                                          2001WO-US00664.
2001WO-US00665.
2001WO-US00666.
2001WO-US00667.
                                                                                                                                              28-JAN-2002; 2002EP-0001165
                                                                                                                                                                                                                                              2001US-0864761
2001US-0328205
        23-DEC-2002 (first entry)
                                                                                                                                                                                                                                                                                                                 WPI; 2002-684061/74.
                                                                                                                                                                                                                                                                           (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention
                                                                                                                                                                                                      30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                      10-OCT-2001;
                                                                                                       EP1239051-A2
                                                                                     Homo sapiens
                                                                                                                                                                          30-JAN-2001;
                                                                                                                                                                                              30-JAN-2001;
                                                                                                                                                                                                                                               23-MAY-2001;
                                                                                                                            11-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the
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Antisense compound which modulates human sentrin expression, useful for treating diseases associated with sentrin expression -

Example 15; Column 38; 29pp; English.

The present invention describes an antisense compound (I) 8-30 nucleotides long targeted to a nucleic acid molecule encoding human sentrin. The antisense compound comprises a phosphorothioate antisense oligonucleotide which inhibits expression of human sentrin. (I) is useful for inhibiting expression of sentrin in human cells or tissues in vitro, for treating humans or other animals suspected of having or being prone to a disease associated with sentrin expression. (I) can also be used for research or diagnostic purposes. The present sequence represents a human sentrin phosphorothioate antisense

Sequence 18 BP; 4 A; 5 C; 3 G; 6 T; 0 other; oligonucleotide from the present invention.

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids. (SL, ABBR3999), a sequence having 65% sequence identity to (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GPPsass as well as downstream components of the signal transduction pathway. (I) is useful of or identifying a specific binding partner. (I) and nucleic acids (II) encoding (I) are useful for diagnosing, monitoring disagnes and treating caused by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating creates the sequence is that of a scanning oligonucleotide useful in examples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to Derwent
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Gaps
                                                                                                                                                                                                                                   Human; epilepsy; chromosome 2; SCN1A; SCN2A; SCN3A; identification;
diagnosis; mutation; chromosome 2q23-q31; neurological disorder;
                           ;
 0.8%; Score 10.6; DB 1; Length 18; 76.5%; Pred. No. 6.8e+02;
                          4; Indels
                                                                                                                                                                                                         Human SCN1A PCR-SSCP PCR primer SEQ ID NO:125.
                         0; Mismatches
                                                1077 GGCTCTTCAGTGAGTGT 1093
                                                                                                                                     BP.
                                                                        17
           76.5%;
                                                                       1 GGCACTTCAGTAACTTT
                                                                                                                                  AAH55881 standard; DNA; 18
                                                                                                                                                                                   (first entry)
           Local Similarity 76.5
                                                                                                                                                                                   04-SEP-2001
Query Match
                                                                                                                                                          AAH55881;
                                                                                                          RESULT 711
                        Matches
                                                                                                                       AAH55881,
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Gaps

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623 GGGACCAGCTCCAGGAG 639

ð

76.5%;

17

GGGCAGAGCTCCGGGAG

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anticonvulsant; neuroprotective; PCR primer; ss.
             WO200138564-A2
     Homo sapiens.
                            26-NOV-1999;
                                                                                                                                                                                    WO9824913-A2
                                                                                                                                                        24-FEB-1999
                                      Rouleau GA,
                  31-MAY-2001
                                                                                                                                                                                          11-JUN-1998
        Synthetic
                                                                                                                                                   AAV95056:
                                                                                                                                        RESULT 712
                                                                                                                                           AAV95056/
g
                                                                                                                                              à
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à 쉄 The present invention describes a method (M1) of determining an individual's predicting the individual's response to medication. The as well as predicting the individual's response to medication. The method comprises determining the genotype of at least one gene selected from SCN1A, SCN2A or SCN3A, or a DNA variant, equivalent, or mutation which shows a linkage disequilibrium. SCN1A, SCN2A and SCN3A are all codium channel genes located on chromosome 2. The idopathic generalised epilepsy (IGE) gene is more specifically localised on chromosome 2423-431. Compounds identified as modilators of the biological activity epilepsy or other neurological Assorders. They have anticonvulsant and neuroprotective activities. AAH55763 to AAH56164 and AAB99674 to AAH59679 represent SCN1A, SCN2A, and SCN3A converse and proteins given in the exemplification of the present invention. epilepsy Determining a predisposition to epilepsy and/or development of epileps comparises determining the genotype of SCNIA, SCNZA and/or SCN3A, or a DNA variant, equivalent, or mutation which shows a linkage disequilibrium -Cossette P, Ragsdale D; Rochefort D, Example 3; Fig 2; 268pp; English. 24-NOV-2000; 2000WO-CA01404. LaFreniere RG, 99US-0167623 WPI; 2001-355945/37. (UYMC-) UNIV MCGILL,

Sequence 18 BP; 4 A; 6 C; 6 G; 2 T; 0 other;

Gaps . 0 0.8%; Score 10.6; DB 1; Length 18; 76.5%; Pred. No. 6.8e+02; tive 0; Mismatches 4; Indels 729 GGGGCCTGGCTGCCGC 745 Query Match
Best Local Similarity 76.5
Matches 13; Conservative

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17 dadadccreccrecrec

Mouse IL-2 receptor g-chain substrate position 399.

Human, IL-2 receptor g-chain, interleukin 2 receptor gamma chain, hammerhead ribozyme; hairpin ribozyme; substrate; expression; cancer, autoimmune digease; psoriasis; allergy; inflammatory disease; graft rejection; ss.

AAV95056 standard; RNA; 18 BP

(first entry)

97WO-US21748. 02-DEC-1997;

96US-0758306 03-DEC-1996;

(RIBO-) RIBOZYME PHARM INC.

McSwiggen JA, Stinchcomb DT;

WPI; 1998-33332/29

Ribozymes targeted to interleukin 2 - useful for treating e.g. cancer, autoimmune disease and allergies

Claim 4; Page 44; 61pp; English.

The present sequence invention describes ribozymes targeted to modulate the synthesis and/or expression of interleukin (IL)-2R gamma encoded RNA. AAV93889 to AAV94574 represent specifically claimed ribozymes, and AAV9455 to AAV95260 represent specifically claimed substrate sequences from the present invention. The ribozymes can be used for the treatment of, e.g. graft rejection, autoimmune disease, cancer, psoriasis, allergy and other inflammatory conditions. The ribozymes are also used to induce tolerance in a recipient to alloantigen from a donor.

Sequence 18 BP; 3 A; 7 C; 4 G; 4 U; 0 other;

Gaps ; 0 0.8%; Score 10.6; DB 1; Length 18; 76.5%; Pred. No. 6.8e+02; Live 0; Mismatches 4; Indels Local Similarity 76.5 nes 13; Conservative Query Match Best

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AAF91219 standard; DNA; 19 BP AAF91219; RESULT 713 AAF91219

(first entry) 04-MAY-2001

Human multi drug resistance-1 gene related sequence SEQ ID NO: 306,

Human, MDR-1; multi drug resistance-1; drug uptake; disease; cancer; inflammatory disease; neuronal disease; CNS disease; cardiovascular disease; PCR primer; ss.

Homo sapiens

WO200109183-A2.

08-FEB-2001.

28-JUL-2000; 2000WO-EP07314.

30-JUL-1999; 99EP-0114938. 22-FEB-2000; 2000EP-0103361.

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Roots I; Brinkmann U, Hoffmeyer S, Eichelbaum M,

WPI; 2001-159855/16.

New polynucleotide encoding a molecular variant Multi Drug Resistance (MDR)-1 polypeptide is useful for diagnosing and treating diseases associated with abnormal MDR-1 expression or function, e.g. cancer -

Disclosure; Page 140; 154pp; English.

us09904568-3.rng

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RESULT 715
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Matches
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                  AAF91220
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The present invention provides nucleotides encoding molecular variants of the human multi drug resistance-1 (MDR-1) protein. These can be used to identify compounds capable of treating multidrug resistance and sensitivity interfering resulting from polymorphisms in MDR-1, which can lead to difficulties in treating cancer, cardiovascular, neuronal, inflammatory and CNS diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention provides nucleotides encoding molecular variants of the human multi drug resistance-1 (MDR-1) protein. These can be used to identify compounds capable of treating multidrug resistance and sensitivity interfering resulting from polymorphisms in MDR-1, which can lead to difficulties in treating cancer, cardiovascular, neuronal, inflammatory and CNS diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polynucleotide encoding a molecular variant Multi Drug Resistance (MDR)-1 polypeptide is useful for diagnosing and treating diseases associated with abnormal MDR-1 expression or function, e.g. cancer-
                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                          Human; MDR-1; multi drug resistance-1; drug uptake; disease; cancer; inflammatory disease; neuronal disease; CNS disease; cardiovascular disease; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                    Human multi drug resistance-1 gene related sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 10.6; DB 1; Length 19; 76.5%; Pred. No. 7.1e+02; ive 0; Mismatches 4; Indels
                                                                                                              Length 19;
                                                                                                          0.8%; Score 10.6; DB 1; Length 1 76.5%; Pred. No. 7.1e+02; Live 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Roots
                                                                                    Sequence 19 BP; 2 A; 5 C; 9 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 3 A; 9 C; 5 G; 2 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hoffmeyer S, Eichelbaum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 140; 154pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                            GGTACAGCGTGGCCCTG 912
                                                                                                                                                                                    19
                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-JUL-1999; 99EP-0114938.
22-FEB-2000; 2000EP-0103361.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-JUL-2000; 2000WO-EP07314
                                                                                                                                                                                  GGCAGACGGTGGCCCTG
                                                                                                                                                                                                                                             AAF91221 standard; DNA; 19
                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-159855/16.
                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                     WO200109183-A2
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                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                              04-MAY-2001
                                                                                                                                    13;
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                                                                                                                                                            968
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                                                                                                            Query Match
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                                                                                                                        Best Loca
Matches
                                                                                                                                                                                                                      RESULT 714
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The present invention provides nucleotides encoding molecular variants of the human multi drug resistance-1 (MDR-1) protein. These can be used to identify compounds capable of treating multidrug resistance and sensitivity interfering resulting from polymorphisms in MDR-1, which can lead to difficulties in treating cancer, cardiovascular, neuronal, inflammatory and CNS diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New polynucleotide encoding a molecular variant Multi Drug Resistance (MDR)-1 polypeptide is useful for diagnosing and treating diseases associated with abnormal MDR-1 expression or function, e.g. cancer -
                                                                                                                                                                    Human, MDR-1; multi drug resistance-1; drug uptake; disease; cancer; inflammatory disease; neuronal disease; CNS disease; cardiovascular disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, MDR-1; multi drug resistance-1; drug uptake; disease; cancer; inflammatory disease; neuronal disease; CNS disease; cardiovascular disease; PCR primer; ss.
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                                                                                                                             Human multi drug resistance-1 gene related seguence SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 10.6; DB 1; Length 19; llarity 76.5%; Pred. No. 7.1e+02; Conservative 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eichelbaum M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 140; 154pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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BP.
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22-FEB-2000; 2000EP-0103361.
AAF91220 standard; DNA; 19
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ID AAF91222 standard; DNA; 19
                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                         WO200109183-A2
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                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                               28-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Brinkmann U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                   04-MAY-2001
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                                         AAF91220
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Gaps

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GGTACAGCGTGGCCCTG 912

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Conservative

Local Similarity

13;

Matches

17 decadacedradecere 1

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Gaps

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The present sequence is a sequencing primer which is used for
                                                                                                                                                                                   Example 5; Column 25; 26pp; English.
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                      3rinkmann
                                                                                      17
                                                                                                         AAD04572;
                                                                       Query Match
                                                                                                                                                              Lan MS,
                                                                                              RESULT 717
                                                                            Matches
                                                                                                                                Homo
                                                                                                                                                                              and
                                                                                                 AAD04572,
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sequencing the human insulinoma-associated antigen, IA-1 cDNA clone. The IA-1 function as a regulatory factor in islet cell transformation. The IA-1 is used as a tumour marker for diagnosis and identification of insulinoma and neuroendocrine tumours. It is also used for identifying cancers. Correct identification of insulinomas and cancers is possible. The IA-1 fragments may be used to immunise animals for the generation of polyclonal and monoclonal antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is that of a primer used in the PCR amplification of the open reading frame of a cDNA clone (see AAA29150) encoding human phosphatidylethanolamine N-methyltransferase-2 (PEMT-2, see AAY79199). The PCR product was subcloned into mammalian expression vector pCI, and PEMT-2 was expressed in rat hepatoma McAAdle-RH777 cells. The invention relates to novel human PEMT2 polymucleotides and protein (see AAY79199), and to methods of using them in the treatment and diagnosis of liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated nucleic acid molecule encoding phosphatidylethanolamine N-methyltransferase protein used to treat phosphatidylethanolamine N-methyltransferase-associated disorders such as liver cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Phosphatidylethanolamine N-methyltransferase-2; PEMT2; human; liver cancer; hepatoma; antitumour; antiproliferative; therapy; diagnosis; PCR primer; ss.
                                                                                                                                                                                                                           0.8%; Score 10.6; DB 1; Length 19;
76.5%; Pred. No. 7.1e+02;
ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match

0.8%; Score 10.6; DB 1; Length 19;
Best Local Similarity 76.5%; Pred. No. 7.1e+02;
Matches 13; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                               Sequence 19 BP; 5 A; 6 C; 5 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 2 A; 7 C; 7 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RESE ) RESEARCH CORP TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 8; Page 57; 111pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disorders, such as liver cancer.
                                                                                                                                                                                                                                                                                                                  475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP
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                                                                                                                                                                                                                                               76.5%;
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                                                                                                                                                                                                                                                                                                               459 GGTCAGCAGCCTGCAGG
                                                                                                                                                                                                                                                                                                                                                         18 GGTGATCCTCCTGCAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ94157 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human PEMT2 PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Walkey CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-256956/22
                                                                                                                                                                                                                                               Local Similarity
hes 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ94157;
                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides nucleotides encoding molecular variants of the human multi drug resistance-1 (MDR-1) protein. These can be used to identify compounds capable of treating multidrug resistance and sensitivity interfering resulting from polymorphisms in MDR-1, which can lead to difficulties in treating cancer, cardiovascular, neuronal, inflammatory and CNS diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel insulinoma-associated neuroendocrine tumor-associated cDNA, useful for diagnosing and identifying insulinoma, neuroendocrine tumors
                                                                                                                                                                                                                                                                             New polynucleotide encoding a molecular variant Multi Drug Resistance (MDR)-1 polypeptide is useful for diagnosing and treating diseases associated with abnormal MDR-1 expression or function, e.g. cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 10.6; DB 1; Length 19;
llarity 76.5%; Pred. No. 7.1e+02;
Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4; Indels
                                                                                                                                                                                                  Roots I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 3 A; 8 C; 5 G; 2 T; 1 other;
                                                                                                                                                                                               Hoffmeyer S, Eichelbaum M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 140; 154pp; English.
                                                                                                                                                  (EPID-) EPIDAUROS BIOTECHNOLOGIE AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  896 GGTACAGCGTGGCCCTG 912
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                                                                                    30-JUL-1999; 99EP-0114938.
22-FEB-2000; 2000EP-0103361.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94US-0246489
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Notkins AL;
                                                                                                                                                                                                                                         WPI; 2001-159855/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-299371/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
les 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-JUN-1992;
08-FEB-2001
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Gaps

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404 CCCGGCTACTAGGGGAC 420

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PD-ATP-binding cassette; PD-ABC; chromosome 19p13.3; spleen; thymus; ds;

form 2 DNA exon 11 3' splice site.

Human PD-ABC

(first entry)

peripheral blood leukocyte; bone marrow; lymph node; dyslipidaemia; cardiovascular disorder; inflammatory disorder; abnormal calcium flux; epilepsy; coronary artery disease; Tangler's disease; atherosclerosis; familial high-density lipoprotein deficiency; fatty liver disease; atherosclerosis, diabetes; insulin resistance; obesity; drug screening; alcoholism; retinal degeneration; hypertension; vascular disease.

CCCGGCTGCTGGGCTAC 18

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The sequence represents a splice site within a DNA molecule encoding human PD-Arp-binding cassette (PD-ABC) protein. PD-ABC maps to chromosome 19P13.3 and is expressed in various tissues including spleen, thymus, peripheral blood leukocytes, bone marrow and lymph nodes. The PD-ABC DNA molecules and proteins are used to diagnose and treat cardiovascular disorders, inflammatory disorders, dysliptdemia, epileps, diseases related to abnormal calcium flux, coronary artery disease, Tangier's disease, fantlial high-density lipoprotein deficiency, atherosclerosis, diabetes, fatty liver disease, insulin resistance, obesity, alcoholism, retinal degeneration, hypertension and vascular disease. The sequences are also used in drug screening assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ^{\prime} Human PD-ABC DNA molecules and proteins for diagnosis and treatment dyslipidaemia, epilepsy and diseases related to abnormal calcium flux
                                                                                                                                                                              PD-ATP-binding cassette; PD-ABC; chromosome 19p13.3; spleen; thymus; ds; peripheral blood leukocyte; bone marrow; lymph node; dyslipidaemia; cardiovascular disorder; inflammatory disorder; abnormal calcium flux; epilepsy; coronary artery disease; Tangier's disease; atherosclerosis; familial high-density lipoprotein deficiency; fatty liver disease; atherosclerosis; diabetes; insulin resistance; obesity; drug screening; alcoholism; retinal degeneration; hypertension; vascular disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 10.6; DB 1;
Pred. No. 7.3e+02;
                                                                                                                                             Human PD-ABC form 1 DNA exon 11 3' splice site.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 37; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wang M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-JAN-2001; 2001WO-US02191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000US-0177889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JUN-2000; 2000US-0215405
                                   AAS08740 standard; DNA; 20
                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (WARN ) WARNER LAMBERT CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tafuri SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-442259/47.
                                                                                                                                                                                                                                                                                                                                                                         WO200153490-A1
                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-JAN-2000;
                                                                                                         26-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                            26-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Johns MA,
                                                                      AAS08740;
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                 AAS08740
RESULT
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New Human PD-ABC DNA molecules and proteins for diagnosis and treatment of dyslipidaemia, epilepsy and diseases related to abnormal calcium flux  $\,$ 

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Wang

Tafuri SR,

Johns MA,

WPI; 2001-442259/47.

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(WARN ) WARNER LAMBERT

23-JAN-2001; 2001WO-US02191. 24-JAN-2000; 2000US-0177889. 30-JUN-2000; 2000US-0215405.

WO200153490-A1. Homo sapiens.

26-JUL-2001

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The sequence represents a splice site within a DNA molecule encoding human PD-ATP-binding cassette (PD-ABC) protein. PD-ABC maps to chromosome 19pl3.3 and is expressed in various tissues including spleen, thymus, peripheral blood leukcytes, bone marrow and lymph nodes. The PD-ABC DNA molecules and proteins are used to diagnose and treat cardiovascular disorders, inflammatory disorders, dyslipidaemia, epilepsy, diseases related to abnormal calcium flux, coronary artery disease, Tangier's disease, fanty liver disease, intoprotein deficiency, atherosclerosis, diabetes, fatty liver disease, insulin resistance, obesity, alcoholism, retinal degeneration, hypertension and vascular disease. The sequences are also used in drug screening assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Secreted protein, human; PRO protein, neoplastic cell growth; tumour; proliferation; leukaemia; lymphoid malignancy; inflammatory disorder; angiogenic disorder; immunologic disorder; PRO6496; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 10.6; DB 1; Length 20; 6.5%; Pred. No. 7.3e+02; ve 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 3 C; 11 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 39; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1248 GGCCATGTGAGGCCAGG 1264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA91053 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      76.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGACAGGTCAGGCGAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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Best Local S
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Gaps ;

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13; Conservative

Best Local Similarity

Matches

1248 GGCCATGTGAGGCCAGG 1264

13

GGACAGGTCAGGCGAGG

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BP.

AAS08831 standard; DNA; 20

RESULT 720

AAS08831

AAS08831;

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Power M;

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The specification describes a beta-secretase enzyme. The enzyme cleaves beta-amyloid precursor protein to produce beta-amyloid peptide. This enzyme is therefore implicated in the production of amyloid plaque components which accumulate in the brains of individuals afflicted with a Latenance's disease. Inhibitors of beta-secretase are administered to a mammalian subject e.g. with Alzheimer's disease or Alzheimer's disease-like pathology to test if they maintain or improve cognitive ability or reduce the plaque burden. The compounds are used for the treatment of amyloidogenic diseases e.g. Alzheimer's disease. PCR primers AAA59530-49 were used to amplify DNA encoding beta-secretase
                                                                                                                                                                                                                                                                                                                                                                                   Purified beta-secretase protein used in assays to discover inhibitors which can be used for the treatment of amyloidogenic diseases e.g. Alzheimer's disease -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.8%; Score 10.6; DB 1; Length 21; Best Local Similarity 76.5%; Pred. No. 7.4e+02; Matches 13; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                     Doane MT, Frigon N, John V,
Tung J, Wang S, McConlogue L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Triple helix forming methylphosphonate oligomer 2104.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Methylphosphonate; MP; triple helix; translation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 3 A; 9 C; 7 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; Page 66; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   575 AGCAGGCCCTCCGTCTG 591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
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                                                                                                                                        10-FEB-2000; 2000WO-US03819.
                                                                                                                                                                                                    99US-0139172.
                                                                                                                                                                                   99US-0119571
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(first entry)
                                                                                                                                                                                                                                          (ELAN-) ELAN PHARM INC.
                                                                                                                                                                                                                                                                                     Basi G,
                                                                                                                                                                                                                                                                Anderson JP, Bab. -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oligonucleoside; ss.
                                                                                                                                                                                                                                                                                                                                              WPI; 2000-533011/48.
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                                                         WO200047618-A2
                                                                                                                                                                              10-FEB-1999;
15-JUN-1999;
                   Homo sapiens
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16-FEB-1995
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                                                                                                  17-AUG-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents a PCR primer used to isolate DNA encoding human PRO5800 protein of the invention. The PRO proteins are secreted proteins. Antagonists or antibodies of PRO polypeptides are useful for diagnosing and treating tumours are associated with gene amplification. Hooplastic cell growth and proliferation in mammals, and those conditions contactived by oversynession and/or activation of the amplified genes. Such conditions include benign or malignant tumours (e.g. renal, liver, kidney, bladder, breast, gastric, ovarian, colorectal, prostate, tidney, bladder, breast, gastric, ovarian, colorectal, prostate, giloblastomas and various head and neck tumours); leukaemias and lymphoid malignancies; neuronal, glial, astrocytal, hypothalamic, and other alignancies; neuronal, glial, stromal and blastocoelic disorders; and inflammatory, angiogenic and immunologic disorders. These may further be used to qualitatively or quantitatively detect the expression of proteins encoded by the amplified genes, and in tumour diagnostics or proparation of a medicament in the treatment of a condition, which is responsive to the PRO polypeptide, its antagonist or anti-PRO antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antibodies against PRO polypeptides, useful for diagnosing and treating tumours are associated with gene amplification, neoplastic cell growth and proliferation in mammals
                                                                                                                                                                                                                                                                                                                                                                                                            Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Beta-secretase; beta-amyloid precursor protein; beta-amyloid peptide; amyloid plaque component; Alzheimer's disease; amyloidogenic disease; inhibitor; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                          Watanabe CK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 10.6; DB 1; Length 20; larity 76.5%; Pred. No. 7.3e+02; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                        Smith V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 5 A; 6 C; 4 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                        Gurney AL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 11; Page 95; 143pp; English.
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                                                                                                                                  99US-0138385.
99US-0144790.
99US-0146843.
99US-0148188.
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99US-0149396.
99US-0150114.
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                                                                                                                                                                                                                                                                                                                                                                                                      Botstein DA, Goddard A,
                                                                                                                                                                                                                                                                                                                                                                (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-071075/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
              WO200075317-A2
                                                                                             15-MAY-2000;
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                                                     14-DEC-2000
                                                                                                                                                         20-JUL-1999;
                                                                                                                                                                              03-AUG-1999
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                                                                                                                                                                                                                                                                                                    -AUG-1999
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RESULT 722

Matches

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Gaps

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ACA07666;

(first entry) 03-JUN-2003

NFKB sub-unit modulating zinzyme substrate #65.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; d-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; postate cancer; colorectal cancer; brain cancer; cescophageal cancer; stomach cancer; brain cancer; cancer; parcreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; cervical cancer; head and neck cancer; ovarian cancer; melanoma; chemotherapy; paclitarel; docetasel; cisplatin; methotrexate; cyclophosphamide; docetasel; cisplatin; methotrexate; cyclophosphamide; docorubin; fluorouracil carboplatin; edatesate; cyclophosphamide; docorubin; fluorouracil carboplatin; edatesate; rheumatoid arthritis; restenosis; crohi s disease; obsity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection;

Homo sapiens.

US2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-0864785

94US-0291932. 92US-0987132. 94US-0245466. 96US-0777916. 15-AUG-1994; 07-DEC-1992; 18-MAY-1994; 23-DEC-1996;

STINCHCOMB D T. STIN/)

MCSW/)

MCSWIGGEN J. DRAPER K G. (DRAP/)

Draper KG; Mcswiggen J, Stinchcomb DT,

WPI; 2003-340953/32

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases

Claim 3; Page 38; 72pp; English

regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (1) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (1) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, cervical, head and neck, overian cancer, melahoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug chemotherapy including paciltaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, caid molecules are also useful for treating inflammatory disease such as acid molecules (2) and artistense nucleic contemporal arthritis, restensing inflammatory disease such as contemporated arthritis, restensing inflammatory disease such as contemporated arthritis, restensing a lunian (2) chomic arthritis restensing and inflammatory disease such as chamatoid arthritis, restensing and property contemporated arthritis, restensing and property contemporate and antistensing and property contemporated arthritis, restensing and property contemporated arthritis, restensing and property contemporated arthritis and property contemporated and property contemporated arthritis and property contemporated and property contemporated arthritish contemporated and p invention describes an enzymatic nucleic acid molecule (I) which down rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft (central nervous system (CNS) and myocardial), glomerulonephritis, sepsits, allergic airway inflammation, inflammatory bowel disease or enzymatic nucleic acid molecule.

Sequence 17 BP; 3 A; 9 C; 4 G; 1 U; 0 other; g

Gaps ·. Length 17; Indels 1; Score 10.4; DB 1; Pred. No. 7e+02; '0; Mismatches 1; 0.8%; 11; Conservative Query Match Best Local Similarity Matches

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730 GGGCCTGCCTG 741 à

17 dececcidectic 6

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ACA06320/c RESULT

BP. ACA06320 standard; RNA; 17

ACA06320;

(first entry) 03-JUN-2003

NFKB sub-unit modulating inozyme substrate #139.

G-Cleaver, amberzyme; cancer; REL-A activity; breatt cancer; human; oesophageal cancer; REL-A activity; breatt cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; human; oesophageal cancer; stomath cancer; lander cancer; pancreatic cancer; cervical cancer; had and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cipplatin; methotrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;

Homo sapiens.

US2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-0864785.

94US-0291932. 92US-0987132. 15-AUG-1994;

94US-0245466. 96US-0777916. 07-DEC-1992; 18-MAY-1994; 23-DEC-1996;

STINCHCOMB D T. (/NILS

DRAPER K G. MCSWIGGEN (DRAP/)

Draper KG; Mcswiggen J, Stinchcomb DT,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases

Claim 3; Page 29; 72pp; English.

regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (1) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzyment nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (1) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, invention describes an enzymatic nucleic acid molecule (I) which down 

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us09904568-3.rng

cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or therapies such as monoclohal antibodies, REL-A-specific inhibitors or cherotherapy including pacilitaxel docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edarrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edarrexate, accid molecules are also useful for treating inflammatory disease nucleic rheumatoid arthritis, restenosis, asthma, crohn's disease, diabetes, rejection, autoimmune disease, lupus, multiple sclerosis, transplant/graft (central nervous system (CNS) and myocardial), glomernlonephritis, sepsis, allergic airway inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule.

Sequence 17 BP; 3 A; 8 C; 4 G; 2 U; 0 other;

Gaps .. 0 Length 17; 1; Indels 0.8%; Score 10.4; DB 1; 91.7%; Pred. No. 7e+02; iive 0; Mismatches 1; Query Match Best Local Similarity 91,7° Matches 11; Conservative

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à g RESULT 728

ACA06587 standard, RNA, 17 BP. ACA06587,

ACA06587;

03-JUN-2003 (first entry)

NFKB sub-unit modulating inozyme substrate #406.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; decleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; postate cancer; colorectal cancer; brain cancer; oesophageal cancer; stomach cancer; brain cancer; pancreatic cancer; cervical cancer; stomach cancer; varian cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; chemotherapy; paclitary resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitary fluorouracil carboplatin; edatrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; rheumatoid arthritis; restendats; Crohn's disease; asthma; diabetes; rheumatoid arthritis; restendats; Crohn's disease; obsity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection;

Homo sapiens.

US2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-0864785 .5-AUG-1994; 

94US-0291932. 92US-0987132. 94US-0245466. 96US-0777916. 07-DEC-1992; 18-MAY-1994; 23-DEC-1996;

STINCHCOMB D T. MCSWIGGEN J. DRAPER K G. (/NILS) (DRAP/) MCSW/

Draper KG; Stinchcomb DT, Mcswiggen J,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression

regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG-2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, ossophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, caid and antisense nucleic acid molecules are also useful for treating inflammatory disease such as rhemmatoric as the useful for treating inflammatory disease such as rhemmatoric as the useful for treating inflammatory disease such as rhemmatoric as the page of the contact of the cancer of an antisense and an arrespondent or reductions. invention describes an enzymatic nucleic acid molecule (I) which down of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases Claim 3; Page 33; 72pp; English. 

Sequence 17 BP; 2 A; 6 C; 5 G; 4 U; 0 other;

Gaps ; 0 0.8%; Score 10.4; DB 1; Length 17; 91.7%; Pred. No. 7e+02; ive 0; Mismatches 1; Indels 11; Conservative

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TGCAGCTGCAGC 4 15

ACA08920,

ACA08920 standard; RNA; 17 BP. (first entry) 03-JUN-2003 ACA08920; 

NFKB sub-unit modulating amberzyme substrate #83.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; protate cancer; colorectal cancer; brain cancer; nesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; cervical cancer; head and neck cancer; ovarian cancer; melanoma; chemotherapy; aloma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; ovachophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; cancer allergic airway inflammation; inflammatory bowel disease; infection;

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-0864785.

rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule.

Query Match 0.8%; Best Local Similarity 91.7%; 827 IGCAGCIGAAGC 838 Matches a à

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The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG^2-+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, crostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, colorectal, brain, oesophageal, stomach, bladder, pancreatic, crovical, head and neck, ovarian cancer, melanoma, lymphoma, glloma or multidrug resistant cancer. The method involves use of other drug therapise such as monoclonal antibodies, REL-A-specific inhibitors or cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, colomential are also useful for treating inflammatory disease such as caid molecules are also useful for treating inflammatory disease such as colomethin, encouracil carboplatin, cerification, gene therapy applications, schmenial repetivation injury cefetival nervous system (CNS) and myocardial), glomerulonophritis, septiminal repetical injury cefetival nervous system (CNS) and myocardial), glomerulonophritis, septiminal any cefetival nervous system control inflammatory bowel disease or marwaric minial aria and any control inflammatory bowel disease or conference represents the substrate of a novel
                                                                                                                                                                                                                                                                                                                                   Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases
                                                                                                                                                                                                                                             Draper KG;
                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3; Page 51; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   enzymatic nucleic acid molecule.
                      94US-0291932.
92US-0987132.
94US-0245466.
                                                                                                                                                                                                                                           Mcswiggen J,
                                                                                             96US-0777916
                                                                                                                                        STINCHCOMB D
                                                                                                                                                                 MCSWIGGEN J.
DRAPER K G.
                                                                                                                                                                                                                                                                                        WPI; 2003-340953/32.
                                                                                                                                                                                                                                        Stinchcomb DT,
                    15-AUG-1994;
07-DEC-1992;
18-MAY-1994;
                                                                                             23-DEC-1996;
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                                                                                                                                                                                        (DRAP/)
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Sequence 17 BP; 3 A; 8 C; 4 G; 2 U; 0 other; Query Match Best Local Similarity Matches

730 GGGGCCTGGCTG 741 14 GGGGCCTGCCTG 3 à

AAS45551 standard; DNA; 20 BP. AAS45551; RESULT 730 AAS45551 

Tumour-specific IgV region H chain, PCR primer gamma. (first entry)

18-DEC-2001

Human; B cell lymphoma; cytostatic; immunostimulator; self-antigen; tumour-specific vaccine; tumour; polyclonal immune response; idiotype-specific anti-lymphoma immune response; PCR primer; ss.

Homo

WO200168682-A1

13-OCT-2000; 2000WO-US28362 20-SEP-2001 

10-MAR-2000; 2000US-0522900

(LARG-) LARGE SCALE BIOLOGY CORP. MCCORMICK A A. TUSE D. (TUSE/) MCCO/

Turpen TH; Reinl SJ,

WPI; 2001-596903/67.

Novel polypeptide vaccine produced in plants, useful for inducing an immune response to a self-antigen on the surface of certain tumour cells

Disclosure, Page 30; 89pp; English.

the function of the state of th The invention relates to a novel polypeptide self-antigen (I) useful as

Sequence 20 BP; 4 A; 8 C; 6 G; 2 T; 0 other;

.; 0.8%; Score 10.4; DB 1; Length 20; 70.0%; Pred. No. 7.7e+02; Indels 0; Mismatches 70.08; Local Similarity 70.0 es 14; Conservative Query Match Matches

0;

Gaps

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Gaps

.; 0

ABZ76936 standard; DNA; 20 RESULT 731 ABZ76936

(first entry) 07-MAY-2003

ABZ76936;

Bovine DGAT BAC-DNA sequencing primer #9.

chromosome 14; Acyl CoA:diacylglycerol transferase; DGAT; enzyme; bovine; milk; meat marbling; low fat; polymorphic; single nucleotide polymorphism; PCR primer; ss.

Bos taurus 

Synthetic

WO2003004630-A2

WO2003004630-A2

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New nucleic acid molecule comprising a sequence of an allele of a polymorphic bovine acyl CoA-diacylglycerol transferase gene useful for testing a mammal for its predisposition for fat content of milk and for
                                                   (ARBE-) ARBEITSGEMEINSCHAFT DEUT RINDERZUECHTER.
                                                                                                                                     Example 1; Page 35; 91pp; English.
               05-JUL-2002; 2002WO-EP07520.
                              06-JUL-2001; 2001EP-0116412,
13-MAX-2002; 2002US-379412P,
                                                                                  WPI; 2003-239205/23.
                                                                   Winter A;
                                                                                                                      meat marbling
16-JAN-2003
                                                                   Fries H,
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The present invention describes a nucleic acid molecule (NA) (I) encoding a bovine acyl CoA-diacylglycerol transferase (DGAT) contributing to or indicative for low fat content of milk and to low meat marbling (intramuscular fat content). Human DGAT is located to chromosome 8, and bovine DGAT is located to chromosome 8, and the mammal for its predisposition for fat content of milk and/or its bovine DGAT is predisposition for meat marbling. The method comprises analysing the redisposition for meat marbling. The method comprises analysing the predisposition for meat marbling. The method comprises analysing the gene encoding DGAT for nucleotide polymorphisms (SNPs) which are connected with the predisposition. The nucleotide polymorphisms are located in the coding region of the DGAT gene and result in substitution, deletion and/or addition of an amino acid sequence of the polypeptide which is encoded by the gene. The nucleotide acid molecule has at the position 10433 and 10434 of the DGAT gene a guanine, 11030 a guanine, 11048 a cytosine or thymine and 11093 a corresponding to position 10433 and 10434 of the DGAT gene two adenine corresponding to position 10433 and 10434 of the DGAT gene two adenine cresidons which is responsible for the regulation for high content of milk and high meat marbling. The nucleotide polymorphisms are located in a region which is responsible for the regulation of the expression of the product of the gene encoding DGAT. ABZ7045 at ABZ7045 and ABBP6035 to invention.

Sequence 20 BP; 7 A; 7 C; 5 G; 1 T; 0 other;

Gaps ·, 0.8%; Score 10.4; DB 1; Length 20; 91.7%; Pred. No. 7.7e+02; tive 0; Mismatches 1; Indels Best Local Similarity 91.7 Matches 11; Conservative Query Match

AGGCCCCCACAG 12

ABZ77002; ABZ77002 

(first entry) 07-MAY-2003

Bovine DGAT PCR primer #38.

866 AGGTCCCCACAG 877 à

ABZ77002 standard; DNA; 20 BP. RESULT 732

Acyl CoA:diacylglycerol transferase; DGAT; enzyme; chromosome 14; bovine; milk; meat marbling; low fat; polymorphic; SNP; single nucleotide polymorphism; PCR primer; ss.

Bos taurus. Synthetic.

New nucleic acid molecule comprising a sequence of an allele of a polymorphic bovine acyl CoA-diacylglycerol transferase gene useful for testing a mammal for its predisposition for fat content of milk and for (ARBE-) ARBEITSGEMEINSCHAFT DEUT RINDERZUECHTER 05-JUL-2002; 2002WO-EP07520. 06-JUL-2001; 2001EP-0116412. 13-MAY-2002; 2002US-379412P. WPI; 2003-239205/23. Fries H, Winter A; 16-JAN-2003

Example 1; Page 36; 91pp; English.

meat marbling

The present invention describes a nucleic acid molecule (NA) (I) encoding a bovine acyl CoA-diacy1glycerol transferase (DGAT) contributing to or indiactive for low fat content of milk and to low meat marbling (intramuscular fat content). Human DGAT is located to chromosome 8, and (intramuscular fat content). Human DGAT is located to chromosome 8, and bovine DGAT is located to chromosome 14. (I) is useful for testing a mammal for its predisposition for fat content of milk and/or its predisposition for meat marbling. The method comprises analysing the gredisposition for medial polymorphisms (SNPs) which are connected with the predisposition. The nucleotide polymorphisms (SNPs) which are connected with the predisposition. The nucleotide polymorphisms are located in the coding region of the DGAT gene and result in substitution, deletion and/or addition of an amino acid sequence of the polypeptide which is encoded by the gene. The coding regular and a cytosine residue, at position 10433 and 10434 of the DGAT gene a guanine, and a cytosine residue, at position 1993 and 1993 a cytosine or thymine, which correlate with a predisposition for low fat content of milk and low meat marbling. The nucleic acid molecule has at the position corresponding to position 10433 and 10434 of the DGAT gene two adenine residues which correlate with a predisposition for high content of milk and high meat marbling. The nucleotide polymorphisms are located in a region which is responsible for the regulation of the expression of the present invention invention.

Sequence 20 BP; 7 A; 7 C; 5 G; 1 T; 0 other;

0; Gaps 0.8%; Score 10.4; DB 1; Length 20; 91.7%; Pred. No. 7.7e+02; tive 0; Mismatches 1; Indels 11; Conservative Local Similarity Query Match Matches Best

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AAZ71860 standard; DNA; 20 BP. (first entry) 10-SEP-2001 AAZ71860; AAZ71860 

RESULT 733

Human biallelic marker upstream amplification primer SEQ ID NO:6216.

Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; single mucleotide polymorphism; SNP; PCR primer; diagnosis; ss.

06-APR-2000; 2000WO-US09054

99US-0288461

08-APR-1999; SISI (-SISI)

PHARM INC

WPI; 2000-619223/59

Karras JG;

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Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                       AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ740 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diseascing methods, as well as the characterisation of the differential efficacious responses to and side
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           effects from pharmaceutical agents acting on a disease as well as other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N.B. The SEQ ID NOs 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 5 A; 2 C; 9 G; 4 T; 0 other;
                                                                                                                                                                                                                                                   Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                            Claim 9; Page 1556; 2745pp; English.
                                                                                                                       99WO-IB00822,
                                                                                                                                                           98US-0082614
98US-0109732
                                                                                                                                                                                                                                                 Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                           map of the human genome
                                                                                                                                                                                                                                                                                    WPI; 2000-013267/01.
                                                                                                                                                                                                               (GEST ) GENSET
                    Homo sapiens.
                                                    WO9954500-A2
                                                                                                                       21-APR-1999;
                                                                                                                                                                            23-NOV-1998;
                                                                                                                                                           21-APR-1998;
                                                                                      28-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                treatment.
                                                                                                                                                                                                                                                   Cohen D,
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The present invention describes an antisense compound (1), 8 to 30 nucleobases in length, that is targeted to a nucleic acid molecule encoding STAT3 (Signal Tranaducer and Activator of Transcription) and which inhibits the expression of it. (1) has antinflammatory, it is used to inhibiting the expression of STAT3 in cells or tissues, treating an animal having a disease or condition associated with STAT3 or a number having a disease or condition daracterised by a reduction in a poptiosis, and inducing apoptosis in a cell. Diseases or conditions that are treated are rheumatoid arthritis, cancer of the breast, prostate, brain, head and/or neck, lenksemia, myeloma, melanoma or lymphoma. (1) can also be used for diagnostic methods in detecting and determining the role of STAT3 in various cell functions, physiological processes and conditions and for diagnostic methods in detecting as an immunostimulator. (1) is used in sandwich and colourimetric assays, involving enzyme conjugation and radiolabeling and is used in diagnostic kits. AAC93150 encodes human STAT3 and AAC93231 encodes mouse STAT3 as given in the exemplification of the present invention. AAC93151 to AAC93230 and AAC93232 to AAC93299 represents a mismatch control.

New antisense compound for inhibiting the expression of signal transducer and activator of transcription 3 (STAT3) in cells or tissues and treating diseases or condition associated with STAT3, such as rheumatoid arthritis and cancer -

Example 2; Page 46; 104pp; English

Gaps , / Match 0.8%; Score 10.4; DB 1; Length 20; Local Similarity 91.7%; Pred. No. 7.7e+02; nes 11; Conservative 0; Mismatches 1; Indels Query Match Matches ð

16 5 GGAAGTTGAGGT

g

AAC93165 standard; DNA; 20 AAC93165; RESULT 734 AAC93165 

BP.

(first entry) 15-FEB-2001

Human STAT3 phosphorothioate antisense oligonucleotide SEQ ID NO:16.

Human; mouse, STAT3; phosphorothioate; antisense oligonucleotide; modulation; signal transducer and activator of transcription; DNA-binding protein; signal transduction; inhibition; apoptosis; inflammatory disease, cancer; antiinflammatory; antirheumatic; cytostatic; immunostimulatory; rheumatoid arthritis; leukaemia; myeloma; melanoma; lymphoma; diagnosis; ss.

sapiens. Homo

WO200061602-A1

19-OCT-2000.

Sequence 20 BP; 4 A; 3 C; 8 G; 5 T; 0 other; 11; Conservative Best Local Similarity Query Match Matches ö 6 GGCAGTTGAGGT 17

823 CTGATGCAGCTG 834 9 CTGATGGAGCTG à

20

; 0

Gaps

.; 0

Length 20;

Score 10.4; DB 1; Length 20; Pred. No. 7.7e+02; n: Mismatches 1; Indels

0;

0.8%;

91.7%;

(first entry) 26-FEB-2002 AAS96782;

AAS96782 standard; DNA; 20 BP.

RESULT 73 AAS96782

antisense phosphorothioate oligodeoxynucleotide #15. Human STAT3

STAT3; human; signal transducer and activator of transcription; ss; STAT; attisense gene therapy; Fas-mediated apoptosis; inflammatory disease; autoimmune disease; rheumatoid arthritis; cancer; breast; prostate; head; neck; brain; leukaemia; myeloma; melanoma; lymphoma; apoptosis; antiinflammatory; immunosuppressive; antirheumatic; antiarthritic; cytostatic

Homo sapiens. 

Synthetic.

US2001029250-A1.

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The invention relates to antisense compounds targeted to a nucleic acid molecule encoding a signal transducer and activator of transcription (STAT) protein, specifically STAT3, where the antisense compounds inhibit the expression of STAT3. The antisense sequences are useful for inhibiting the expression of STAT3 in cells or tissues, inducing are apoptosis in cells, and sensitising cells to apoptosis. They are also useful for treating an animal having a disease or condition associated with STAT3. These disorders include inflammatory or autoimmuse the presst, prostate, brain and head and neck and leukaemias, myslomas. They breast, prostate, brain and head and neck and leukaemias, myslomas. They are sequences of the invention can disease or conditions apoptotic signals. The sequences of the invention can be used in clinical research, for detecting and determining the role of STAT3 in various cell suscitions and physiological processes and for diagnosing conditions and associated with the expression of STAT3. The sequences represent cDNA and also associated with the expression of STAT3. The sequences represent cDNA and also associated with the expression of STAT3.
                                                                                                                                                                                                                                                        Novel antisense compound useful for treating and diagnosing inflammatory diseases and cancers, is targeted to a nucleic acid molecule encoding signal transducer and activator of transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KSHV; gamma herpes virus; glycoprotein B; vaccine; infection;
human Kaposi's sarcoma-associated herpes virus; probe; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 10.4; DB 1; Length 20;
11.7%; Pred. No. 7.7e+02;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 encoding human STAT3 and human STAT3 oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KSHV DNA polymerase antisense oligonucleotide HVLQB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 3 C; 8 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 13; 21pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT84695 standard; DNA; 21 BP
                                                                                                           06-APR-2000; 2000WO-US09054.
                                                      11-JAN-2001; 2001US-0758881.
                                                                                            99US-0288461
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                                                                                                                                                                                                                         WPI; 2002-009991/01
                                                                                                                                                (KARR/) KARRAS J G.
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                                                                                            08-APR-1999;
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                   11-OCT-2001
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Gaps

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Gaps
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                                                                                                     DNA encoding glycoprotein B of retroperitoneal fibromatosis and Kaposi's sarcoma associated herpes viruses - useful in vaccines for treatment of herpes infection or for detection of viral DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Herpes virus DNA polymerase and corresponding nucleotide sequence used in the detection and treatment of herpes virus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Retroperitoneal fibromatosis herpes virus, detection, infection, Kaposi's sarcoma herpes virus; viral DNA, viral RNA, vaccine;
                                                                                                                                                                                           Claimed type 3 oligonucleotides (AAT84694-96) are specific non-degenerate oligonucleotides for the human Kaposi's sarcomasssociated herpes virus (KSHV) DNA polymerase (gB). They can be used for detecting, amplifying or characterising KSHV polymucleotides encoding DNA polymerase (see AAT84697).
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                                                                                                                                                                                                                                                                                                                   0.8%; Score 10.4; DB 1; Length 21; 31.7%; Pred. No. 7.8e+02; Ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KSHV DNA polymerase specific oligonucleotide HVLQB.
                                                                                                                                                                                                                                                                                       Sequence 21 BP; 3 A; 4 C; 11 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Todaro GJ;
                                                                                                                                                                Claim 37; Page 76; 138pp; English.
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                                            Strand K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT51587 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                   91.78;
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95US-0001148
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                                                                                                                                                                                                                                                                                                                                                                              522 CCTGCCGGAGGA 533
                                                                                                                                                                                                                                                                                                                                  Local Similarity 91.7 ses 11; Conservative
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            (UNIW ) UNIV WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antigen; antibody; ss.
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                                          Rose IM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-132644/12
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14-JUL-1995;
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                                         Bosch ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                                                                                                                                                                                                                                     Query Match
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Matches
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Antisense therapy; antiproliferative; antiinflammatory;

Antisense oligonucleotide #10.

(first entry)

30-MAR-2001

AAF45161;

BP.

161/c AAF45161 standard; RNA; 15

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                                                                                                                                                                                                                                                                                                                                                                                                                         SpA domain D; Ig binding region; gamma chain; B-cell superantigen; sAg; superantigen, heavy chain variable region; VH3 restricted antibody; VH; protein-A; VH2G; combinatorial library; B-lymphocyte; vaccine; DNA probe; hybridization; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A B-cell superantigen (8Ag) is a fragment of SpA D domain that specifically binds the Fab portion of variable region restricted antibodies. The sAg is used to enhance production of VH, especially VH3, restricted Abs. To detect Ig gamma chain expression, the antisense sequence given in AAQ64857 was used as probe. Detection of VH families used the sense oligonucleotides given in AAQ64858-60. (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                       Gaps
family. They may also be used in the design and screening of anti-viral drugs. Antibodies raised against the polymerase or fragments of it, may be used in the detection of herpes virus infection and for drug targetting for the therapy of herpes virus
                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stimulating prodn. of variable region gene family restricted antibodies - through B-cell super-antigen vaccination
                                                                                                                       Length 21
                                                                                                                                                     Indels
                                                                                                                      Score 10.4; DB 1;
Pred. No. 7.8e+02;
0; Mismatches 1;
                                                                                        Sequence 21 BP; 3 A; 4 C; 11 G; 3 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 24; 130pp; English.
                                                                                                                                                                                                                                                                                                                                                                                             Ig gamma chain probe gamma-CHI.
                                                                                                                                                                                                                                                                                        BP.
                                                                                                                    Ouery Match
Best Local Similarity 91.7%;
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92US-0969936
                                                                                                                                                                                                                                                                                     AAQ64857 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                (updated)
(first entry)
                                                                                                                                                                              522 CCTGCCGGAGGA 533
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                                                                                                                                                                                                            CCTGCTGGAGGA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9409B18-A1
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18-0CT-1994
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                                                           infection
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                                                                                                                                                                                                                                                        RESULT 738
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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an ancisense oligonacleotide, (for Insulin-11ke Growth Factor [IGF]-1 receptor. [Debinding protein [IGFB]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is one such antisense oligonucleotide. The method is useful for ameliorating the effects of psoriasis, idhthyoais, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                             cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin discorder; Insulin-like Growth Factor I receptor; IGFP-1; pityriasis; IGF binding protein; IGFB-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hypermeovascular condition; hyperplasia; kidney disease; neovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic aid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 15 BP; 4 A; 5 C; 6 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Edmondson SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MURD-) MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 15; Page 115; 201pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  21-JUN-2000; 2000WO-AU00693.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-0140345.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wraight CJ, Werther GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              729 GGGGCCTGGCTGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-041421/05.
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                                                                                                                                                                                                                                                                                                                                                              WO200078341-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-JUN-1999;
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AAF49864
ID AAF4986
XX
AC AAF4986
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Gaps

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0.8%; Score 10.4; DB 1; Length 23; 70.0%; Pred. No. 8e+02; ive 0; Mismatches 6; Indels

704 TCCTTGATTCTGTGGCCCAG 723 TCCTTGACCAGGCAGCCCAG 21

à

Local Similarity 70.0

Best Loca Matches

Query Match

(first entry) 30-MAR-2001 IGF-I oligonucleotide #824

Antisense therapy, antiproliferative, antiinflammatory, antipsoriatic, cytostatic, dermatological, cardiant, virucide, ophthalmological; keloid, skin disorder, Insulin-like Growth Factor I receptor; IGF-1, pityriasis; IGF binding protein, IGFB-2, IGFBF3, inflammation, psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba, keratosis; neoplasia; scleroderma, wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia, kidney disease; neoblation of the retina; ss.

Homo sapiens

WO200078341-A1.

28-DEC-2000

21-JUN-2000; 2000WO-AU00693

99US-0140345. 21-JUN-1999;

(MURD-) MURDOCH CHILDRENS RES INST,

Werther GA, Edmondson SR; Wraight CJ,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation -

Example 8; Page 66; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligomucleotide, (for Insulin-IRe Growth Factor [167]-1 receptor, 1GF binding protein [1678]-2 or 1GFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligomucleotide which can be used to design the antisense oligomucleotide which can be used to design the antisense oligomucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serbornhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the settin, a hypernevescular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia.

Sequence 15 BP; 0 A; 6 C; 5 G; 4 T; 0 other;

Gaps ·, 0.8%; Score 10.2; DB 1; Length 15; 0.0%; Pred. No. 6.8e+02; ve. 0; Mismatches 3; Indels 80.08; Query Match 0.8 Best Local Similarity 80.0 Matches 12; Conservative

o;

729 GGGGGCCTGGCTGCC 743 Grerecerecerede 15

à Д

ACA06585 standard; RNA; 17 BP RESULT 741 ACA06585

ACA06585;

03-JUN-2003 XXXXXXXXX

NFKB sub-unit modulating inozyme substrate #404.

(first entry)

G-cleaver, amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; cervical cancer; prostate cancer; colorectal cancer; pancreatic cancer; cervical cancer; pancreatic cancer; cervical cancer; pancreatic cancer; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemcherapy; paclitear!, docetaxel; cisplatin; methotrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restences; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;

Homo sapiens

US2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-0864785.

Ь. MCSWIGGEN (MCSM/ 

Draper KG; Stinchcomb DT,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases

Claim 3; Page 33; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (1) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRM), where (I) is an inozyme, zinzyme, q-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a partient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2, The enzymatic and anticident cation, especially Mg^2, The enzymatic and anticident cation, especially Mg^2, The enzymatic and unitidivity resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, gencitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule.

Sequence 17 BP; 3 A; 6 C; 6 G; 2 U; 0 other;

Gaps ò 0.8%; Score 10.2; DB 1; Length 17; 80.0%; Pred. No. 7.5e+02; ve 0; Mismatches 3; Indels 80.08; Conservative Best Local Similarity Matches 12; Conserva Query Match

0

3

5-AUG-1994;

07-DEC-1992; 18-MAY-1994; 23-DEC-1996;

94US-0291932. 92US-0987132. 94US-0245466. 96US-0777916.

(/NILS)

STINCHCOME D T. DRAPER K G. (DRAP/)

Mcswiggen J,

us09904568-3.rng

715 GTGGCCCAGCAGCAG 729

2 daddcccudcudcad 16

BP. AAX67028 standard; RNA; 18

AAX67028;

(first entry) 20-JUL-1999

Mouse B7 hairpin ribozyme target SEQ ID NO:3660.

Arthritic condition, graft tolerance, immune response, target, cleavage, hammerhead ribozyme, hairpin ribozyme, human; rabbit, mouse, collagenase, stromelysin; synovial membrane, joint, arthritis, osteoarthritis, rheumatoid arthritis; autoimmune disease, allergy; inflammation; diagnosis; ss

Mus sp.

WO9618736-A2.

20-JUN-1996

22-NOV-1995;

95WO-US15516

95US-0541365 05-OCT-1995

95US-0390850. 95US-0426124. 95US-0432874. 95US-0434509. 94US-0354920. 94US-0363253. 94US-0363254. 23-DEC-1994; 23-DEC-1994; 17-FEB-1995; 20-APR-1995

95US-0000951. 95US-0512861 07-JUL-1995 02-MAY-1995 04-MAY-1995

(RIBO-) RIBOZYME PHARM INC.

DT; Stinchcomb Draper K, Gustofson J, McSwiggen J, Pavco P, Stinchcon Beigelman L, Karpeisky A, Modak A, Usman N, Burgin A; Matulic-Adamic J, Jarvis T, Thompson JD, Wincott F; Draper K, Gustofson J, McSwiggen J,

WPI; 1996-300653/30.

Enzymatic nucleic acid molecules having a hammer-head motif - used for the treatment of arthritis, induction of graft tolerance or treatment of auto-immune diseases 

Claim 10; Page 215; 307pp; English.

The present invention describes a novel enzymatic nucleic acid (ENA) having a hammerhead motif (HM) comprising; (i) at least 5 ribose residues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification. The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for enhancing graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly specific. The present sequence is used in the exemplification of the

Gaps ., Length 18; Indels 0.8%; Score 10.2; DB 1; 53.3%; Pred. No. 7.7e+02; iive 4; Mismatches 3; Sequence 18 BP; 1 A; 4 C; 4 G; 9 U; 0 other; Conservative Local Similarity ., Query Match Best Loca Matches X S

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1032 TGCAGCTGACTCTTC 1046 vecuecueauveuc ò ద

RESULT 743

BP AAV57794 standard; DNA; 18

AAV57794;

18-NOV-1998

(first entry)

Human chromosome 18 PCR mapping primer clone 47r.

Manic-depressive illness, susceptibility, genotype, diagnosis, chromosomal marker, polymorphic marker; chromosome 18; human; myo-inositol monophosphatase protein; IMP-18p; PCR primer; ss.

Homo sapiens. Synthetic

WO9818963-A1

07-MAY-1998.

97WO-US19381 28-OCT-1997;

(USSH ) US DEPT HEALTH & HUMAN SERVICES. 96US-0029278 28-OCT-1996;

Esterling LE; Berrettini WH, Detera-Wadleigh SD, Be Goldin LR, Sanders AR, Yoshikawa T; Badner JA, Gershon ES,

WPI; 1998-272247/24.

New isolated IMP.18p myo-inositol monophosphatase - used to develop products for determining susceptibility to manic depressive illness and as targets for preventive and therapeutic treatments

Example 5; Page 71; 118pp; English.

A method has been developed for determining a genotype associated with increased susceptibility to manic-depressive (MD) illness. The method comprises determining the genotype of an affected individual with at least one polymorphic marker localised within the chromosomal region defined by and including markers D185843 and D185869 and determining the genotype associated with increased susceptibility to MD disorder. The method can be used for determining susceptibility to MD illness including bipolar disorder, genetic counselling of individuals from families affected with MD illness, and aid in the differential diagnosis of MD illness from other psychiatric pathologies. Products from the present invention can also be used to obtain modulators of IMP.18p myoinositol monophosphatase protein activity and as targets for preventive and therapeutic treatments. The present sequence represents a PCR primer used in the mapping of human chromosome 18 for determining the genotype of MD illness susceptibility, used in an example from the present invention 

Sequence 18 BP; 2 A; 4 C; 4 G; 8 T; 0 other;

Gaps .. 0 0.0%; Score 10.2; DB 1; Length 18; 10.0%; Pred. No. 7.7e+02; ve 0; Mismatches 3; Indels 80.08; Query Match 0.8 Best Local Similarity 80.0 Matches 12; Conservative

643

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The present sequence is one of a number of oligonucleotides designed to target different regions of the human NF-kappa-B p65 subunit, which is a member of the Rel/NF-kappa-B family of transcription factors.

Rel/NF-kappa-B proteins are involved in a diverse set of signaling pathways involving stress, apoptosis, cancer, growth, infection and inflammation. Antisense oligonucleotides are able to inhibit expression of the p65 subunit and may therefore be used in the treatment of disorders associated with NF-kappa-B p65 subunit expression. They may be used as a prophylaxis to prevent or delay infection, inflammation or tumour formation. Antisense compounds may also be used for research and
                                                                                                                                                                                                                                Human NF-kappa-B p65 subunit antisense oligodeoxynucleotide ISIS# 23738.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense compounds which inhibit the expression of the human NF-Kappa-B p65 subunit (p65) useful for treating diseases associated with p65 expression and as prophylaxis to prevent of delay infection, inflammation or tumor formation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diagnostics because they hybridise to nucleic acids encoding NF-kappa-B p65 subunit. The effect of antisense oligonucleotides on NF-kappa-B p65 subunit mRNA levels was measured using real-time quantitative PCR and Northern blot analysis. Antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oligonucleotides were synthesised on an automated DNA synthesiser.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   phosphorothioate and optionally some sugars may be 2' methoxyethyl"
                                                                                                                                                                                                                                                                  Human; anti-inflammatory; cytostatic; antimicrobial; infection; antisense inhibition; inflammation; transcription factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                some internucleoside bonds are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8%; Score 10.2; DB 1; Length 18; Best Local Similarity 80.0%; Pred. No. 7.7e+02; Matches 12; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 2 A; 6 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 15; Column 40; 33pp; English.
                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monia BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                    or
                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0199859.
                                                                                                                        AAA27086 standard; DNA; 18
                                   15 ÁGCTACÁGAAGCACT 1
                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cowsert LM,
629 AGCTCCAGGAGCTCT
                                                                                                                                                                                                                                                                                                                                                                                                1..18
/*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                    /note=
                                                                                                                                                                                                                                                                                                           apoptosis; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-410858/35.
                                                                                                                                                                                                                                                                                                                                                                               Key
modified_base
                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-NOV-1998;
                                                                                                                                                                                                21-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US6069008-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-MAY-2000
                                                                                                                                                              AAA27086;
                                                                                                           AAA27086/c
                                                                                        RESULT
                                                                                                                            ð
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Primer 1 and 2 (AAQ86978-79) were used to amplify a target nucleotide sequence from methicillin resistant S. aureus (MRSA). The target is detected with a probe specifically derived from the mecA gene of S. aureus and S. epidermidis. The mecA gene product has no homology with known PBPs (penicillin-binding proteins). The new probes allow for the rapid identification of all MRS, eradicating need for labour intensive in vitro cultivation and physiological assays.
                                                                                                                                                                             MRSA; methicillin resistant Staphylococcus aureus; probe; hybridisation; mecA; MRSE; Staphylococcus epidermis; primer; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detection of methicillin resistant Staphylococcus - using an oligo:nucleotide derived from the mecA gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seguence 18 BP; 7 A; 1 C; 4 G; 6 T; 0 other;
                                                                                                                                                       Primer 1 to amplify MRSA target DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Page 11; 14pp; German.
                                                                                                                                                                                                                                                                                                                              93DE-4338119.
                                                                                                                                                                                                                                                                                                                                                          93DE-4338119
                                                                      AAQ86978 standard; DNA; 18
CTACACAGGACCAGG 1
                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                  Springer W;
                                                                                                                                                                                                                                            Staphylococcus aureus.
                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1995-180108/24.
                                                                                                                                                                                                                                                                                                                                                                                       ÅĞ.
                                                                                                                                                                                                                                                                                                                                                                                       (FARB ) BAYER
                                                                                                                                                                                                                                                                                                                                                                                                                  Endermann R,
                                                                                                                             17-JAN-1996
                                                                                                                                                                                                                                                                                                                               08-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                          08-NOV-1993;
                                                                                                                                                                                                                                                                       DE4338119-A1
                                                                                                                                                                                                                                                                                                  11-MAY-1995.
15
                                                                                                  AAQ86978;
                                        RESULT 745
                                                        AAQ86978
                                                                                  g
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Non-immunogenic; epitope; T-cell; immunogenicity; immune system; SK; immunoglobulin; therapeutic; streptokinase; de-immunised; 708; De-immunised 708 Vk constructing flanking primer DIVK7. BP. 441 AAAGTIGCTGAAGTT 455 AAAGATGATGCAGTT 16 AAV81061 standard; DNA; 18 (first entry) 03-MAR-1999 AAV81061; 7 RESULT 746 AAV81061/ à a

ö

Gaps

0

Length 18; 3; Indels

0.8%; Score 10.2; DB 1; 80.0%; Pred. No. 7.7e+02;

0; Mismatches

Best Local Similarity 80.0 Matches 12; Conservative

Query Match

primer; ss. Synthetic

ö

Gaps

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à

421 CTAGACAGGAGCACG 435

1

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Reducing immunogenicity of proteins - by modifying the amino acid sequence of the protein to eliminate potential epitopes for T-cells of a given species
                                                                                                                                       Example 3; Fig 16; 77pp; English.
                                                                         (BIOV-) BIOVATION LTD
                                                                                                  WPI; 1999-045301/04.
WO9852976-A1
                        21-MAY-1998;
                                                             02-DEC-1997;
                                          21-MAY-1997;
                                    14-APR-1998;
                                                31-JUL-1997;
28-NOV-1997;
            26-NOV-1998
                                                                                                                                                                                                                                     vh and vk
                                                                                      Carr FJ;
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97GB-0010480. 97GB-0016197. 97GB-0025270.

97US-0067235

98WO-GB01473. 98GB-0007751.

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The invention relates to a method for the production of non-immunogenic proteins. The method comprises determining at least part of the amino acid sequence of the protein, (b) identifying in the amino acid sequence one or more potential epitopes for T-cells (T-cell epitopes) of the given species, and (c) modifying the amino acid sequence to eliminate at least one of the T-cell epitopes identified in step (b) thereby to eliminate or reduce the immunogenicity of the protein when exposed to the limmune system of the given species. A method of analysing a pre-existing protein to predict the basis for immunogenic responses is also provided. The methods can be used particularly for reducing the immunogenicity of immunoglobulins or therapeutic proteins. e.g. Streptokinase (SX). The products can be used for diagnosts and therapy. Sequences AAV81047-68 represent oligonucleotides used for the construction of de-immunised 708
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 2 A; 5 C; 4 G; 7 T; 0 other;
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RESULT 748

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ó
                                    Gaps
                                    o,
 0.8%; Score 10.2; DB 1; Length 1
80.0%; Pred. No. 7.7e+02;
live 0; Mismatches 3; Indels
Query Match 0.85
Best Local Similarity 80.07
Matches 12; Conservative
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1026 CCAAAGTGCAGCTGA 1040 ccaagcragaacraa 3 17 ò d

AAV51978 standard; DNA; 19 AAV51978; RESULT 747 AAV51978/ THE STANDARD STANDARD

02-FEB-1999 (first entry)

Zea mays genome reverse PCR primer #274.

Polymorphic marker; allele-specific; probe; amplification; PCR primer; hybridisation; plant; hybrid certification; genetic contribution; progeny; back-cross; hybrid; ancestry; corn; ss.

Synthetic Zea mays.

WO9824796-A1

11-JUN-1998

97WO-US21782. 01-DEC-1997;

```
AAV51705-V52008 are reverse PCR primers used to amplify fragments of th Zea mays genome in order to detect polymorphic markers. Such markers can be used in the construction of allele-specific primers and probes for amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, to monitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a back-crossed plant with an ancestral plant.
                                                                                                                                                                           species allele-specific oligonuclectide probes and primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8%; Score 10.2; DB 1; Length 19; Best Local Similarity 80.0%; Pred. No. 8e+02; Matches 12; Conservative 0; Mismatches 3; Indels
                                                                                              Sapolsky
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 5 A; 6 C; 7 G; 1 T; 0 other;
                                                                                                                                                                                                                                     Example 1; Page Page 54; 65pp; English.
                                                                                                Murigneux A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               729 GGGGCCTGGCTGCC 743
                96US-0032069.
                                                                                                                                                                                              - useful for plant breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15 ecerecerecerece 1
                                                        (AFFY-) AFFYMETRIX INC
                                                                                              Lemieux B,
                                                                                                                                     WPI; 1998-333252/29
07-MAR-1997;
                02-DEC-1996;
                                                                                              Landry BS,
                                                                                                                                                                           Brassica
```

can

; 0

Gaps

Polymorphic marker; allele-specific; probe; amplification; PCR primer; hybridisation; plant; hybrid certification; genetic contribution; progeny; back-cross; hybrid; ancestry; corn; ss. Zea mays genome reverse PCR primer #275. ВЪ AAV51979 standard; DNA; 19 (first entry) 02-FEB-1999 AAV51979; AAV51979/ 

Synthetic.

Zea mays.

WO9824796-A1.

11-JUN-1998

97WO-US21782. 01-DEC-1997; 97US-0813507. 96US-0032069 02-DEC-1996; 07-MAR-1997:

(AFFY-) AFFYMETRIX INC

Sapolsky RJ; Murigneux A, Landry BS, Lemieux B,

WPI; 1998-333252/29

Brassica species allele-specific oligonucleotide probes and primers for plant breeding useful

Example 1; Page Page 54; 65pp; English.

AAV51705-V52008 are reverse PCR primers used to amplify fragments of the Zea mays genome in order to detect polymorphic markers. Such markers can

; 0

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amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, to monitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a back-crossed plant with an ancestral plant.
in the construction of allele-specific primers and probes for
                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                           Human; steroidogenesis; acute regulatory protein; hStAR; analysis; mutation; detection; prenatal; genetic defect; congenital; protein; lipoid adrenal hyperplasia; treatment; prevention; gene; replacement therapy; hypercholesterolaemia; primer; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         steroidogenesis acute regulatory protein (hStAR) cDNA. The hStAR gene can be analysed for mutations to detect (e.g. prenatally) genetic defects associated with congenital lipoid adrenal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated human steroidogenesis acute regulatory protein gene - . for detection of mutation(s) of this gene that cause congenital lipoid adrenal hyperplasia
                                                                                                                                                              .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is a PCR primer (nt 717-738) for the human
                                                                                                                                                                                                                                                                                                                                                                                 Steroidogenesis acute regulatory protein antisense PCR primer
                                                                                                                                 Length 19;
                                                                                                                            0.8%; Score 10.2; DB 1; Length 1
10.0%; Pred. No. 8e+02;
ve 0; Mismatches 3; Indels
                                                                                                  Seguence 19 BP; 5 A; 6 C; 7 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 7; Page 4; 89pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Strauss JF;
                                                                                                                                                                                                                                                                                         AAT39478 standard; DNA; 20 BP
                                                                                                                                            80.08;
                                                                                                                                                                                        729 GGGGGCCTGGCTGCC 743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          96WO-US03896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95US-0410540
                                                                                                                                                                                                                 dcerectrectrect 1
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                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                             12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Miller WL,
                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9629338-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-MAR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                    21-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                    15
                                                                                                                                                                                                                                                                                                                       AAT39478
nseq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lin D,
                                                                                                                                                          Matches
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This sequence represents a primer for the coding sequence for the alpha epithelial sodium channel a (alphaENACa). AlphaENACa (see AAW34529) and alphaENACA (see AAW34530) represent the sequences of the invention. The two sodium channels are alternatively spliced forms of the amiloride-sensitive salt channel alpha subunit and can be used in membrane-transport or binding assays to identify substances that enhance or block perception of a salty taste. Enhances could be used as salt substitutes and blockers could be used to mask salty tastes in foods and
                                                                                                                                                                                                                                                                         Alpha epithelial sodium channel, alphaENACa, alphaENACb; binding assay, amiloride-sensitive salt channel alpha subunit, membrane-transport; salt substitute, salty taste blocker; PCR primer; amplify; ss.
                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ively spliced epithelial sodium channel alpha subunit - useful in screening assays for salty taste enhancers or
                               ..
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0
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                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 10.2; DB 1;
80.0%; Pred. No. 8.1e+02;
 Score 10.2; DB 1;
Pred. No. 8.1e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                             Primer alphaEN-S2 for alphaENAC coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 5 C; 7 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Column 9; 33pp; English.
                            0;
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0
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                                                         1246 GTGGCCATGTGAGGC 1260
0.8%;
ilarity 80.0%;
Conservative
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94US-0202654.
                                                                                   16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGCTCCTGGGGCTAT 18
                                                                                                                                                            AAT99084 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYJO) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                    (first entry)
                                                                             GTGGCCATGCCAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      629 AGCTCCAGGAGCTCT
                                                                                                                                                                                                                                                                            Alpha epithelial sod
amiloride-sensitive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ľi X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-031814/03
 Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
nes 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pharmaceuticals.
                                                                                                                                                                                                                                                                                                                                                  Rattus rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alternatively
                                                                                                                                                                                                                                                                                                                                                                                                                                        23-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blackshaw S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-FEB-1994;
                                                                                                                                                                                                                    24-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                              US5693756-A.
                                                                                                                                                                                                                                                                                                                                                                                                           02-DEC-1997
                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               blockers
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Matches
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8, 2004, 16:40:18

Search completed: January Job time : 26 secs

probe to isolate a 1.6 kb insert, including an ORF for a 285 residue protein. When it was cloned into pSPORT and expressed in COS-1 cells cotransfected with pP450scc abd pADX, it increased the level of pregnenolone synthesis from cholesterol or

Sequence 20 BP; 4 A; 7 C; 6 G; 3 T; 0 other;

20-alpha-hydroxycholesterol

hyperplasia (CAH), or its transmission to children. CAH can be treated by protein or gene replacement therapy, which can also be used to prevent or treat hypercholesterolaemia.

A human adrenal cortex CDNA library was screened with a mouse StAR

GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.

OM nucleic - nucleic search, using sw model

January 8, 2004, 16:43:28; Search time 13 Seconds (without alignments) 1.953 Million cell updates/sec Run on:

us-09-904-568-3 1355 1 gggcaggcagttgaggtgga.....gtgtttcaggcaggcccgg 1355 Title: Perfect score: Sequence:

IDENTITY NUC Gapop 10.0 , Gapext 0.5 Scoring table:

541 seqs, 9368 residues Searched:

1082 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 12 Maximum DB seq length: 50

Post-Processing: Minimum Match 0% Maximum Match 100% Listing first 550 summaries

rni3.seg:\* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	Sequence 65, Appl Sequence 16, Appl Sequence 16, Appl Sequence 16, Appl Sequence 16, Appl Sequence 22, Appl Sequence 23, Appl Sequence 29, Appl Sequence 25, Appl Sequence 19, Appl Sequence 140, Appl Sequence 18, Appl Sequence 33, Appl Sequence 33, Appl Sequence 33, Appl Sequence 33, Appl Sequence 34, Appl Sequence 35, Appl Sequence 35, Appl Sequence 35, Appl Sequence 352, Appl Sequenc	,
SUMMARIES	US-09-302-681-65 US-09-667-135-7 US-08-67-135-7 US-08-788-6228-16 US-08-788-6218-16 US-08-997-22 US-08-644-67-22 US-09-244-7945-29 US-09-244-796-29 US-09-247-190-29 US-09-247-190-29 US-09-247-190-29 US-09-247-190-29 US-09-247-190-29 US-09-247-190-29 US-09-325-6031-2 US-09-325-6031-2 US-09-422-978-11451 US-09-422-978-11451 US-09-422-978-11451 US-09-422-978-11451 US-09-422-978-11451 US-09-422-978-11451 US-09-422-978-11451 US-09-422-978-11451 US-09-422-978-11451 US-09-422-978-11451 US-09-422-978-11451 US-09-422-978-11451 US-09-461-753-55 US-09-40-43-33 US-09-470-443-33 US-09-470-443-33 US-09-470-443-33 US-09-470-443-35 US-09-470-443-35 US-09-470-443-35 US-09-470-443-35 US-09-470-443-35 US-09-470-443-35	
Length DB	22222222222222222222222222222222222222	
% Query Match		
Score		
Result No.	0000 00 00 000000 000 000 00 000000 000 000000	

6	Sequence 30, Appl Sequence 49, Appl	equence 49,	equence 137	guence 137,	squence 43,	quence 253	equence 25	quence 1	quence 26	equence 3,	quence 11	quence 3,	quence 3,	quence 42	equence 13	quence 11	duence 11	quence 8,	quence 3,	quence 13	quence 48	quence 2,	quence 24	quence 7,	quence 18	quence 41	quence 4,	quence 4,	quence 59	quence 12. quence 16	quence 68	quence 52	quence 12.	quence 47	quence 52	quence 623	quence 11	quence 62 quence 62	quence 6,	quence 3,	quence 14,	quence 2,	quence 2,	quence	quence 2,	quence 14,	Juence 4, Appl Juence 2029, A	quence 4, Ap	quence 4, Appl	quence 17, App	quence 645, Ap	quence 2, Appl mence 684. An	quence 2548	Sequence 33, Appl Sequence 33, Appl
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	.2 0.9 17 1 172-09-371-1607	0.9 17 1 IIS-09-371-772R-1678 Sequence	sequence	2 0.9 I/ I US-US-3/I-7/2B-1679 Sequence	2 0.9 17 1 US-09-371-772B-2697 Seguence	2 0 9 17 1 TR-09-371-77-84	appropriate the control of the contr	2 0.9 1/ 1 US-09-3/1-1/28-3069 Sequence	.2 0.9 17 1 US-09-371-772B-3213 Sequence	.2 0.9 17 1 US-09-371-772B-3387 Sequence	.2 0.9 17 1 US-09-371-772B-3660 Semience	.2 0.9 17 1 HS-09-371-772R-4161	.2 0.9 17 1 TELOGENET CALL	and an	Sequence	.z 0.9 I/ 1 US-09-3/1-772B-4722 Sequence	.2 0.9 17 1 US-09-371-772B-5116 Seguence	.2 0.9 17 1 HS-09-371-772B-EE79	2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	sequence	.2 0.3 1/1 US-09-3/1-7/2B-6439 Sequence	.2 0.9 17 1 US-09-371-772B-6624 Segience	.2 0.9 17 1 US-09-371-7728-6701	2 0 9 17 1 DCH TIONE CONTRACTOR	Sequence	2 0.3 1/ 1 FCI-US96-U9641-17 Sequence	2 0.9 18 1 US-08-584-040-3044 Segience	2 0.9 18 1 IIS-09-371-772B-1472	2 12 1 115 - 10 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	color to the color	0.9 13 1 US-08-242-664-14 Sequence	0.9 13 1 US-08-484-138-14 Segmence	0.9 13 1 PCT-US95-06379-14	0 0 14 1 110-10 14 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	control of the control of the sequence	0.9 14 1 US-08-683-839B-15 Seguence	0.9 14 1 US-08-674-168-10 Segmente	0 9 14 1 115 00 000 000 000 000 000 000 000 00	0.3 14 1 US-U8-846-UZIA-14 Sequence	0.9 15 1 US-08-365-189-10 Sequence	0.9 15 1 US-08-208-886C-29	0 9 15 1 115 00 174 174 0	Sequence	IS I US-08-469-557-29 Sequence	. 0.9 15 1 US-08-290-793B-29 Sequence	: 0.9 15 1 US-08-606-505B-62	0 9 15 1 110 00 115 1 00 00 00 00 00 00 00 00 00 00 00 00 0	C. T. T. US-US-IIIS-446-3 Sequence	: 0.9 15 1 US-09-177-359-26 Seguence	0.9 15 1 US-09-616-990-62	0.9 15 1 HS-08-R12-951B-1	0 9 15 1 115 00 012 012 013 013	or the transfer of the sequence	0.9 IS I US-08-747-2 Sequence	0.9 15 1 US-08-784-747-3 Sequence	0.9 15 1 US-09-409-778-9 Sequence	0.9 15 1 US-09-409-778-10	0.9 16 1 HS-0R-232-0873-E	O 9 15 1 110 00 000 CAN D SEQUENCE	0 0 17 1 170 00 170 200 Sequence	2.2 11 1 US-08-158-349 Sequence	0.9 1/ 1 US-09-328-501-14 Sequence	0.9 17 1 US-08-984-709A-45 Sequence	0.9 17 1 US-08-584-040-1844 Segmente	0.9 17 1 HS-08-584-040-7538	0.9 17 1 119-00-537-7370-115	O 14 14 CO OC 14 C	T - CS - C	0.9 1/ 1 US-US-7/7-710A-14 Sequence 14	0.9 17 1 US-09-371-772B-389 Sequence 36	0.9 17 1 US-09-371-772B-4638 Segmence 46	0.9 17 1 PCT-US91-03680-7	1.8 0.9 15 1 IIS-08-041-500-2	Z - C - C - C - C - C - C - C - C - C -	15 1 US-U8-12/-954-50 Sequence 50	1.8 0.9 15 1 US-08-337-025-2 Sequence 2.	1.8 0.9 15 1 US-08-276-099A-8 Semisore 8	1.8 0.9 15 1 US-08-182-9684-201	TR 1	1 to 0 1 to 1 to 0 1 to	15 I US-08-291-932A-378 Sequence 378,	1.8 0.9 15 1 US-08-334-847-570 Sequence 570,	1.8 0.9 15 1 US-08-334-847-606 Sequence 606.	1.8 0.9 15 1 US-08-334-847-631 Semience 631	1.8 0.9 15 1 HS-08-363-2401-142	1.8 0.9 15 1 US-08-363-240A-541 Seminar 11.	1.8 0.9 15 1 IIS-08-343-24015 Common 511,	9 15 1 HG-08-383-240A-858 Sequence 658,	i.e. e.e. 15 i US-US-ISI-BYU-B	

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21 GGTGCGAGCGCAGACTGCGGG 1
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                                       5661
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                                                                      Sequence 3
                   Sequence
Sequence
Sequence
       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Oligonuclectide primer corresponding to NADH US-09-302-681-65
                                                                                                                                                                                                                                                                     APPLICANT: Herrnetedt, Corrina
APPLICANT: Ghosh, Sounitra S.
APPLICANT: Ghosh, Sounitra S.
APPLICANT: Clevenger, William
APPLICANT: Clevenger, William
APPLICANT: Davis, Robert E.
TITLE OF INVENTION: DIANNOSTIC METHOD BASED ON
TITLE OF INVENTION: DIANNOSTIC METHOD BASED ON
TITLE OF INVENTION: ORDATIFICATION OF EXTRAMITOCHONDRIAL DNA
FILE REPERENCE: 660088.416C1
CURRENT APPLICATION NUMBER: US/09/302,681
CURRENT FILING DATE: 1999-04-30
NUMBER OF SOID NOS: 108
SOFTWARE FEBELSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>;</u>
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Fatent No. 6521749
Fatent No. 6521749
FAPLICANT: UNIONATION:
FAPLICANT: Vincent Ling
FILE REFERENCE: GNN-007
FILE REFERENCE: GNN-007
CURRENT APPLICATION NUMBER: US/09/667,135
CURRENT FILING DATE: 2000-09-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 1.6%; Score 22; DB 1; Length 22; Best Local Similarity 100.0%; Pred. No. 2.5; Matches 22; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
 US-09-402-923A-420
US-09-371-772B-5660
US-09-371-772B-561
US-US96-00331-19
US-08-702-105A-33
US-08-702-110A-33
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Best Local Similarity 90.5%; Pred. No. 21;
Matches 19; Conservative 0; Mismatches 2;
                                                                                                                                        ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              34 AGCTACGCAAAATCTTAGCATA 55
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                                                                                                                                                                                                  08-00-302-681-65
; Sequence 65, Application US/09302681
: Patent No. 6441149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ) OTHER INFORMATION: primer US-09-667-135-7
                                                                                                                                                                                                                                        ; Patent No. 6441149
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US-09-667-135-7/c
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11.6
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LENGTH: 22
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                                                                                                                               APPLICANT: Kilburn, Douglas G.
APPLICANT: Miller, Robert C.
APPLICANT: Warren, Richard A.J.
APPLICANT: Gilkes, Neil R.
TITLE OF INVENTION: Polysaccharide binding fusion proteins
TITLE OF INVENTION: and conjugates
NUMBER OF SEQUENCES: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
1.3%; Score 17.6; DB 1; Length 24;
Best Local Similarity 83.3%; Pred. No. 32;
Matches 20; Conservative 0; Mismatches 4; Indels
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CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/249,037C
FILING DATE: 435
PILON APPLICATION DATA: 435
PILON APPLICATION TOWN THE US/08/249,037C
FILING DATE: 08-APR-1994
PILON APPLICATION NUMBER: US/0865,095
PILON APPLICATION NUMBER: US 07/865,095
PILON APPLICATION NUMBER: US 07/603,987
PILING DATE: 25-OCT-1990
PROOF APPLICATION NUMBER: US 07/216,794
PILING DATE: 08-JUL-1988
ATTONNEY/AGBRT INPORMATION:
NAME: KUNG, VIOLA T.
REGISTRATION NUMBER: 41,131
RELECOMMUNICATION INPORMATION:
TELEPHONE: (650)328-4400
                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Rae-Venter Law Group, P.C.
STREET: P.O.Box 60039
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   188 CCGCCCCCCCCGGACGCCGACG 211
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US-08-788-622B-16/c
; Sequence 16, Application US/08788622B
; Patent No. 5962289
; GENERAL INFORMATION:
APPLICANT: Kilburn, Douglas G.
; APPLICANT: Miller, Robert C.
; APPLICANT: Warren, Richard A.U.
RESULT 3
US-08-249-037C-16/c
Sequence 16, Application US/08249037C
Patent No. 5928917
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (650)328-4477
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
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(650)328-4477
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LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20; Conservative
                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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US-08-052-997-22
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APPLICANT: Gilkes, Neil R.
TITLE OF INVENTION: Polysaccharide binding fusion proteins TITLE OF INVENTION: and conjugates
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
1.3%; Score 17.6; DB 1; Length 24;
Best Local Similarity 83.3%; Pred. No. 32;
Matches 20; Conservative 0; Mismatches 4; Indels
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                                                                                                                                                                               COUNTRY: C.A.

COUNTRY: C.A.

ZIP: 94306

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHIN Release #1.0, Version #1.30
SOFTWARE: PATCHIN DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/788,622B
FILING DATE: JANNAT:
APPLICATION NUMBER: US/08/7897
FILING DATE: 24-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/865,095
FILING DATE: 08-APR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/603,987
FILING DATE: 08-APR-1992
PRIOR APPLICATION NUMBER: US 07/603,987
FILING DATE: 08-UUJ-1988
ATTORNEY/AGENT INFORMATION:
NAME: KAUSY, VIOLA T.
REFERENCE/DOCKET NUMBER: CEDT.002.06US
TELEPHONE: (650)328-4400
TELEPHONE: (650)328-4477
INFORMATION POR SEQ ID NO: 16:
SEGUENCE CHARACTERISTICS:
                                                                                                        ADDRESSEE: Rae-Venter Law Group, P.C. STREET: P.O.Box 60039
CITY: Palo Alto
SIATE: CA
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P.O.Box 60039
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Patent No. 612417
GENERAL INFORMATION:
APPLICANT: Miller, Robert C.
APPLICANT: Warren, Richard A.U.
APPLICANT: Gilkes, Neil R.
TITLE OF INVENTION: Polysaccharide by
TITLE OF INVENTION: Polysaccharide by
MUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Rae-Venter Law Group, P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ccarcccarccccarcccarca
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MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: P.O.Box
CITY: Palo Alto
STATE: CA
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US-08-788-621B-16/C
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COMPUTER READALE FORM:

WINDIUM TIPE: RICHER COMMANDER FORM:

APPLICATION NUMBER: US 00/240,037

RICHING DATE: WART 1944

RELIANG DATE: WART 1944

APPLICATION NUMBER: US 00/240,037

RELIANG DATE: WART 1944

RELIANG DATE: WART 1944

APPLICATION NUMBER: US 00/260,997

RICHER APPLICATION DATE: US 00/260,997

RICHER APPLICATION DATE: US 00/260,997

RICHER APPLICATION DATE: US 00/260,997

RICHER APPLICATION NUMBER: US 00/260,997

COUNTY: NEARINGTON DOWNER: US 00/260,997

CONDESSED DOWNERS 00/260,997

COND
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ATTORNEY/AGENT INFORMATION:

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Gaps
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APPLICANT: Roberts, Richard W.
APPLICANT: Roberts, Richard W.
APPLICANT: ADDITION: Richard W.
TITLE OF INVENTION: BELECTION OF PROTEINS USING RNA-PROTEIN
TITLE OF INVENTION: FUSIONS
FILE REFERENCE: 00786/350006
CURRENT APPLICATION WUMBER: US/09/244,794A
CURRENT APPLICATION NUMBER: 60/035,963
PRIOR PILING DATE: 1997-01-27
PRIOR PLING DATE: 1997-01-27
PRIOR PLING DATE: 1997-01-27
PRIOR FILING DATE: 1997-01-27
PRIOR APPLICATION NUMBER: 60/064,491
PRIOR FILING DATE: 1997-01-27
PRIOR APPLICATION NUMBER: 09/007,005
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APPLICANT: Roberts, Richard W.
APPLICANT: Roberts, Richard W.
TAPLICANT: LAW RAHE
TITLE OF INVENTION: FELECTION OF PROTEINS USING RNA-PROTEIN
TITLE OF INVENTION: FUSIONS
TITLE OF INVENTION: FUSIONS
TITLE OF INVENTION: FUSIONS
CURRENT APPLICATION NUMBER: 10/90/007,005B
CURRENT FILING DATE: 1999-01-14
EARLIER APPLICATION NUMBER: 60/035,963
EARLIER PILING DATE: 1997-01-27
EARLIER FILING DATE: 1997-11-06
EARLIER FILING DATE: 1997-11-06
SEQ ID NO 29
SEQ ID NO 29
IENGRARE: FASELSEQ for Windows Version 4.0
                                                                                                                                                               DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.2%; Score 16.4; DB 1; Length 18; 94.4%; Pred. No. 29; Ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                Indels
                                                                                                                                                     Query Match
1.2%; Score 16.8; D
Best Local Similarity 90.0%; Pred, No. 35;
Matches 18; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 29, Application US/09244794A; Patent No. 6214553
FERERAL INFORMATION:
APPLICANT: Szostak, Jack W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 29, Application US/09007005B
Patent No. 6258558
GENERAL INFORMATION:
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                       ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-684-672-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 94.4%;
Matches 17; Conservative
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       : ORGANISM: Homo sapiens
US-09-244-794A-29
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ORGANISM: Homo sapiens
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US-09-244-794A-29
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LENGTH: 18
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APPLICANT: KERE, Juha
APPLICANT: SCHLESSINGER, David
APPLICANT: GLAS CHAFELLE, Albert
APPLICANT: SRIVASTAVA, Anand Kumax
TITLE OF INVENTION: MOLECULAR CLONING OF THE ANHIDROTIC
TITLE OF INVENTION: ECTODERMAL DYSPLASIA GENE
NUMBER OF SEQUENCES: 3
CORRESPONDENCES: 3
CORRESPONDENCES: BURNS, DOANE, SWECKER & MATHIS, L.L.P.
STREET: P.O. BOX 1404
CITY: Alexandria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length 21;
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CITY:

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COMPUTER READABLE FORM:

MEDIUM TYPE:

COMPUTE:

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1.2%; Score 16.8; Dest Local Similarity 90.0%; Pred. No. 35; Matches 18; Conservative 0; Mismatches
      NAME: O'Shaughnessy, Brian P.
REGISTRATION NUMBER: 32,747
REFERENCE/DOCKET NUMBER: 9594/81-2189
TELECOMUNICATION INFORMATION:
TELEPHONE: (202) 289-1200
TELEFAX: (202) 289-674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; CLONE: Yeast artificial chromosome
US-08-052-997-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 22, Application US/08684672
Patent No. 5700926
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                                                                                                                                                                                                                                          22:
                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: unknown
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
                                                                                                                                                            TELEFAX: (202) 289-6674
TELEX: 248516
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORIGINAL SOURCE:
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US-08-684-672-22
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RESULT 12
US-09-238-710-29
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Sequence 29, Application US/09247190

Factor No. 6261804

GENERAL INFORMATION:
APPLICANT: SCORCAK, Jack W.
APPLICANT: Roberts, Richard W.
APPLICANT: Liu, Rihe
ITILE OF INVENTION: FUSIONS
FILE REFERENCE: 00786/35005

CURRENT APLICATION WUMBER: US/09/247,190

CURRENT FILING DATE: 1999-02-09

EARLIER PRILING DATE: 1999-01-21

EARLIER PILING DATE: 1997-01-21

EARLIER PILING DATE: 1997-01-21

EARLIER PILING DATE: 1997-01-21

EARLIER PILING DATE: 1997-01-21

EARLIER PILING DATE: 1999-01-21

SARLIER FILING DATE: 1999-01-21

SARLIER PILING DATE: 1999-01-21

SARLIER PILING DATE: 1998-01-14

NUMBER OF SEQ ID NOS: 38

SOFTWARE: FASTERE FULLOR DATE: 1998-01-14
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Fatent No. 6281344
GENERAL INFORMATION:
APPLICANT: SZOSTAK, U.
APPLICANT: SZOSTAK, U.
APPLICANT: ROBERTS, Richard W.
TITLE OF INVENTION: SELECTION OF PROTEINS USING RNA-PROTEIN
TITLE OF INVENTION: FUSIONS
FILE REPERRER OF 1996
CURRENT APPLICATION NUMBER: 05/09/24,796
CURRENT FILING DATE: 1999-02-05
FARLIER APPLICATION NUMBER: 60/035,963
EARLIER APPLICATION NUMBER: 60/04491
FARLIER PILING DATE: 1997-01-27
FARLIER FILING DATE: 1997-11-06
FARLIER FILING DATE: 1997-11-06
FARLIER FILING DATE: 1998-01-14
NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
                                       1.2%; Score 16.4; DB 1; Length 18; 94.4%; Pred. No. 29; ive 0; Mismatches 1; Indels
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1.2%; Score 16.4; D
Best Local Similarity 94.4%; Pred. No. 29;
Matches 17; Conservative 0; Mismatches
                                                                                                                              679 GIGGIATITGGGAGCCAG 696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          679 GTGGTATTTGGGAGCCAG 696
                                                                                                                                                                1 GTGGTATTTGTGAGCCAG 18
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                                                       Local Similarity 94.4°
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US-09-247-190-29
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CRCANISM: Homo sapiens
US-09-244-796-29
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US-09-244-796-29
US-09-007-005-29
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                                           Query Match
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Matches
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US-09-302-681-76/c
; Sequence 76, Application US/09302681
; Sequence 76, Application US/09302681
; Patent No. 6441149
; GENERAL INFORMATION:
; APPLICANT: Hermsteadt, Corrina
; APPLICANT: Clevenger, William
; APPLICANT: Clevenger, William
; APPLICANT: Pavis, Robert E
; TITLE OF INVENTION: QUANTIFICATION OF EXTRAMITOCHONDRIAL DNA
; TITLE OF INVENTION: QUANTIFICATION OF EXTRAMITOCHONDRIAL DNA
; FILE REFERENCE: 660088-416C1
; CURRENT APPLICATION UNDHER: US/09/302,681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Oligonucleotide primer corresponding to NADH CHER INFORMATION: dehydrogenase encoding mitochondrial DNA US-09-302-681-76
                                                                                                                                                                                                                                                                     Sequence 29, Application US/09238710A
Fatent No. 6518018
GENERAL INFORMATION:
APPLICANT: Szottak, Jack W.
APPLICANT: Szottak, Jack W.
APPLICANT: Szottak, Jack W.
APPLICANT: INFORMATION:
TITLE OF INVENTION: SELECTION OF PROTEINS USING RNA-PROTEIN
TITLE OF INVENTION: FUSIONS
FILE REFERENCE: 00786/350004
CURRENT APPLICATION NUMBER: US/09/238,710A
CURRENT FILING DATE: 1999-01-28
EARLIER APPLICATION NUMBER: 60/035,963
EARLIER FILING DATE: 1997-01-27
EARLIER FILING DATE: 1997-01-27
EARLIER FILING DATE: 1997-01-27
EARLIER FILING DATE: 1998-01-14
NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
             Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 16.4; DB 1; Length 18; Pred. No. 29; 0; Mismatches 1; Indels
                                                          Indels
          DB 1;
Query Match 1.2%; Score 16.4; D
Best Local Similarity 94.4%; Pred. No. 29;
Matches 17; Conservative 0; Mismatches
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 76
LENGTH: 21
                                                                                                    679 GTGGTATTTGGGAGCCAG 696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     679 GIGGIATITGGGAGCCAG 696
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 94.4
Matches 17; Conservative
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; ORGANISM: Homo sapiens
US-09-238-710-29
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DB 1; Length 21;

1.2%; Score 16.2;

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TITLE OF INVENTION: Marek's Disease Virus Genes and Their
TITLE OF INVENTION: Marek's Disease Virus Genes and Their
TITLE OF INVENTION: Use in Vaccines for Protection Against Marek's Disease
NUMBER OF SEQUENCES: 29
ADDRESSEE: Birch, Stewart, Kolasch and Birch, LLP
STREET: P.O. Box 747
CITY: Falls Church
STATE: VA
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CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/709,731A
FILING DATE: 05-JUL-1996
CLASSIETCATION: 424
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US. 08/499,474
FILING DATE: 07-JULY-1995
ATTORNEY/AGENT INFORMATION:

NAME: MUMBER: 28,977
REFERENCE/DOCKET NUMBER: 1644-110FPC
TELEPHONE: (703)-205-8000
TELEPHONE: (703)-205-8000
TELEPHONE: (703)-205-8000
TELEPHONE: (703)-205-8050
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
Best Local Similarity 85.7%; Pred. No. 48;
Matches 18; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.2%; Score 16.2; D
85.7%; Pred. No. 60;
tive 0; Mismatches
                                                                                                                                                                                                                                                                         APPLICANT: Lee, Lucy F
APPLICANT: Nazerian, Keyvan
APPLICANT: Witter, Richard L
APPLICANT: Wu, Ping
APPLICANT: Yanagida, No. 6322780oru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              519 CAACCTGCCGGAGGAGCAGCT 539
                                                                                                                                                                                                                Sequence 5, Application US/08709731A Patent No. 6322780 GENERAL INFORMATION:
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; Sequence 6, Application US/09136959A
; Patent No. 624852.
; GENERAL INFORMATION;
; APPLICANT: HABERHAUSEN, Gerd
                                                                  71
                                                                                        21 GCATACTCTTCAATCAGCCAC 1
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                                                                51 GCATACTCCTCAATTACCCAC
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: sing
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Best Local Similarity
Matches 18; Conservat
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ZIP: 22040-0747
                                                                                                                                                                      RESULT 14
US-08-709-731A-5/c
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GENERAL INFORMATION:
APPLICANT: Weston, Brent W.
APPLICANT: Hiler, Kara B.
TITLE OF INVENTION: Antisense Fucosyltransferase Sequences and Methods of TITLE OF INVENTION: Use Thereof FILE REFERENCE: Weston and Hiller CURRENT APPLICATION NUMBER: US/09/556,031
CURRENT FILING DATE: 1200-04-20
PRIOR FILING DATE: 1999-04-26
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
LIBRIGHH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
APPLICANT: SOBEK, Harald
TITLE OF INVENTION: REDUCTION OF CROSS-CONTAMINATIONS IN NUCLEIC ACID
TITLE OF INVENTION: AMPLIFICATIONS
FILE REFERENCE: 1614-80.65
CURRENT APPLICATION NUMBER: US/09/136,959A
CURRENT FILING DATE: 1998-08-20
PRIOR FILING DATE: 1997-08-20
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 6
LENGTH: 20
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Patent No. 6383809
GENERAL INFORMATION:
APPLICANT: C. FEAN Bennett
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF CYTOHESIN-1 EXPRESSION FILE REFERENCE: RTS-0195
CURRENT APPLICATION NUMBER: US/09/702,246
CURRENT FILING DATE: 2000-10-30
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US-09-556-031-2
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                                                                                                                                                                                                                                                                   TYPE: DNA
CRGANISM: Artificial Sequence
FEATURE:
COTHER INFORMATION: Description of Artificial Sequence:Primer
US-09-136-959A-6
                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
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Pred. No. 52;
0; Mismatches
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1.2%;
Best Local Similarity 89.5%;
Matches 17; Conservative 0
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Best Local Similarity
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Sequence 19, Application US/09322624

Batent No. 6548734

GENERAL INFORMATION:

APPLICANT: Glimcher, L et al.

TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO MODULATION OF TITLE OF INVENTION: CARTILAGE GROWTH BY MODULATION OF NEATH TITLE OF INVENTION: US/09/322,624

CURRENT APPLICATION NUMBER: US/09/322,624

CURRENT FILING DATE: 1999-05-28

MUNBER OF SEQ ID NOS: 20

SOFTWARE: PALENTING DATE: 198-05-28

MUNBER OF SEQ ID NOS: 20

SOFTWARE: PALENTING DATE: 200

LENGTH: 20
                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                       0;
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                                                                                                                                                                                                                                                                                        Query Match
1.2%; Score 15.8; DB 1; Length 20;
Best Local Similarity 89.5%; Pred. No. 52;
Matches 17; Conservative 0; Mismatches 2; Indels
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US-09-702-246-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
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Pred. No. 52
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Best Local Similarity 89.5%;
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: synthetic construct
                                                                                                            NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 19
US-09-422-978-10530/c
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                                                                                                                           SEQ ID NO 25
LENGTH: 20
Thu Jan
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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Cohen, Daniel

APPLICANT: Chumakov, Ilya

APPLICANT: Chumakov, Ilya

TILE OF INVENTION: Biallelic markers for use in constructing a high density...

TILE TO INVENTION BIALLELIC MARKER: 1999-10-20

CURRENT FILING DATE: 1999-04-21

EARLIER APPLICATION NUMBER: US 60/109,732

EARLIER FILING DATE: 1999-11-23

EARLIER FILING DATE: 1999-11-23

EARLIER FILING DATE: 1999-11-23

EARLIER FILING DATE: 1998-11-23

EARLIER FILING DATE: 1998-11-23
; LOCATION: 1..21
; OTHER INFORMATION: downstream amplification primer 99-1333 for SEQ 2665, in comple
US-09-422-978-10530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ) LOCATION: 1..20 ; OTHER INFORMATION: downstream amplification primer 99-6707 for SEQ 3586, in compl US-09-422-978-11451
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APPLICANT: ROSE, TIMOTHY M.
APPLICANT: BOSCH, MARNIX
APPLICANT: STRAND, KURT
APPLICANT: STRAND, KURT
APPLICANT: TODARO, GEORGE J.
TITLE OF INVENTION: DNA POLYMERASE OF GAMMA HERPES VIRUSES
TITLE OF INVENTION: ASSOCIATED WITH KAPOSI'S SARCOMA AND RETROPERITONEAL
TITLE OF INVENTION: FIBROMATOSIS
NUMBER OF SEQUENCES: 152
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                   0; Mismatches
                                                                                                                        1.2%; Score 15.8;
89.5%; Pred. No. 59;
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; Sequence 11451, Application US/09422978
; Patent No. 6537751
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; Patent No. 5925733
                                                                                                                                                                                                                                                    1107 TGTAGTTTTCTGTTTAATT 1125
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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Best Local Similarity 94.1
Matches 16; Conservative
                                                                                                                                                      Best Local Similarity 89.5
Matches 17; Conservative
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California
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US-08-680-326-140/c
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STATE: Ca
COUNTRY:
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                                                                                                                        Query Match
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Patent No. 6537751

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Cohen, Daniel
APPLICANT: Chumakov, Ilya
TITLE OF INVENTION: Biallelic markers for use in constructing a high density...
FILE REFERENCE: GENSET. 020CP1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT PILING DATE: 1999-10-20
FERLIER PILING DATE: 1999-04-21
FERLIER PILING DATE: 1999-10-20
FERLIER PILING DATE: 1998-11-23
FERLIER PILING DATE: 1998-11-23
FERLIER FILING DATE: 1998-11-23
FERLIER FILING DATE: 1998-04-21
FERRIER FILING DATE: 1998-04-21
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ORGANISM: Homo Sapiens FEATURE: NAME/KEY: primer\_bind

SEQ ID NO 10530 LENGTH: 21

Gaps

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Query Match
1.1%; Score 15.4; DB 1; Length 21;
Best Local Similarity 94.1%; Pred. No. 72;
Matches 16; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                             Sequence 89, Application US/08720229
Fatent No. 6022542
GENERAL INFORMATION:
APPLICANT: Bose, Timothy M.
APPLICANT: Bosch, Marnix L.
APPLICANT: Strand, Kurt
TITLE OF INVENTION: GLYCOPROFIN B OF THE RFHU/KSHV
TITLE OF INVENTION: SUBFAMILY OF HERPES VIRUSES
NUMBER OF SEQUENCES: 100
CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CUMINER: U.S.

CUNTER SEADMBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM Compatible
OPERATURG SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/720,229
FILING DATE: 26-SEP-1996
CLASSTFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: SCHIEf, J. Michael
REGISTRATION NUMBER: 40,253
REFERENCE/POCKET UNBER: 2938-20002.00
TRIEDFONE: (415) 813-5600
TELEFRA: (415) 813-5600
TELEFRA: (415) 814-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-07-977-284A-89/c

Sequence 89, Application US/07977284A

Parent No. 5558988

GENERAL INFORMATION:
APPLICANT: Prockop, Darwin J.
APPLICANT: Ala-Kokko, Leena
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Prockop, Darwin J.
Ala-Kokko, Leena
Williams, Charlene J.
                                                                                                                                              569 TGCTCCAGCAGGCCCTC 585
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Baldwin, Clinton
                                                                                                                                                                                   17 rccrccascasscrc 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (412)
                                                                                                                                                                                                                                                                          US-08-720-229-89/c
         US-08-804-439A-89
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-720-229-89
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1.1%; Score 15.4; DB 1; Length 21;
Best Local Similarity 94.1%; Pred. No. 72;
Matches 16; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 22
US-08-804-439A-89/c
; Sequence 89, Application US/08804439A
; Patent No. 601556;
GENERAL INFORMATION:
APPLICANT: Rose, Timothy M.
APPLICANT: Bosch, Marnix L.
APPLICANT: Strand, Kurt
ITLE OF INVENTION: GLYCOPROTEIN B OF THE RFHV/KSHV
ITLES OF INVENTION: SUBFAMILY OF HERPES VIRUSES
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 4225 Executive Square, Ste 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STWIE: CA
COUNTRY: USA
ZIP: 92037

COMPUTER READABLE FORM:
MEDLIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
SURRENT APPLICATION DATA: P6804,439A
FILING DATE: February 21, 1997
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Haile, Lisa A.
REGISTRATION NUMBER: 38,347
PURPERMICH DOCKET NUMBER: 09176/004001
                OPERATING SYSTEM: PC_DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/680,326
                                                                                                                       CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Schiff, U. Michael
REGISTRATION NUMBER: 29936
TELECOMMUNICATION INFORMATION:
TELEFHONE: (415) 494-0792
TELEFAX: (415) 494-0792
TELEFAX: (415) 494-0792
TELEFAX: 706141
INFORMATION FOR SEQ ID NO: 140:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDENRES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   585
IBM PC compatible
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 678-5070
TELEFAX: (619) 678-5099
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LENGTH: 21 base pairs
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 La Jolla
CA
                                                                                                         FILING DATE:
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ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 5948611ris
STREET: One Liberty Place - 46th Floor
CITY: Philadelphia
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APPLICANT: Ala-Kokko, Leena
APPLICANT: Ala-Kokko, Leena
APPLICANT: Williams, Charlene J.
APPLICANT: Ritvaniemi, Pertti
APPLICANT: Baldwin, Clinton
APPLICANT: Abmad, Nilofer Nina
TITLE OF INVENTION: Methods of Detecting A Genetic
NUMBER OF SEQUENCES: 293
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcork
STRREY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.1%; Score 15.2; D
85.0%; Pred. No. 71;
tive 0; Mismatches
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MEDIUM TYPE: DISKETTE, 3.5 INCH
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows 3.1
SOFTWARE: WORDPERFECT 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/256,426B
FILING DATE: 03-FEB-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/10964
FILING DATE: 12-NOV-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/977,284
FILING DATE: 13-NOV-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCT/US93/10964
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REFERENCE/DOCKET NUMBER: TJU-1082
TELECOMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 26
US-08-256-426B-89/c
i Sequence 89, Application US/08256426B
sequence No. 5948611
i GENERAL INFORMATION:
APPLICANT: Prockop, Darwin J.
                         FILING DATE: 23-MAR.1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Neeley, Richard L.
REGISTRATION NUMBER: 30,092
REBERNCE/DOCKET NUMBER: UCAL.
TELEPHONE: 415 853 5070
TELERAX: 415 857 663
TELEX: 380816COLEYPA
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERIES: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs
TYRE: nucleic acid
STRANDENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               548 TGCTGGCAGGCATGCACACA 567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 recreecreecargeccaca 1
                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA (synthetic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 85.09
Matches 17; Conservative
             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Mark DeLuca
                                                                                                                                                                                                                                                                                                                                                              linear
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US-08-410-540-8
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Patent No. 5807678
GENERAL INFORMATION:
APPLICANT: Miller, Walter L.
APPLICANT: Lin, Dong
APPLICANT: Straus III, Jerome F.
TITLE OF INVENTION: ASSOCIATED WITH CONGENITAL LIPOID ADRENAL HYPERPLASIA
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooley Godward Castro Huddleson & Tatum
STREET: P S Palo Alto Square
CITY: P AD ALTO
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
APPLICANT: Hopkinson, Ian
APPLICANT: Ahmad, Nilofer Nina
TITLE OF INVENTION: METHODS OF DETECTING A GENETIC
TITLE OF INVENTION: MEDISPOSITION FOR OSTEOARTHRITIS
NUMBER OF SEQUENCES: 261
CORRESPONDENCE ADDRESS:
ADDRESSES: Moodcock, Washburn, Kurtz, Mackiewicz & No. 5558988ris
STREET: One Liberty Place, 46th floor
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                              COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: HOPPY disk

COMPUTER: IBM C COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WordPerfect 5.1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/977,284A

FILING DATE:

APPLICATION: 435

PRIOR APPLICATION: 435

PRIOR APPLICATION DATA:

APPLICATION NUMBER:

FILING DATE:

ATTORNEY/AGENT INFORMATION:

NAME: DELUCA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: DeLuca, Mark
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: TJU-0697
TELEPHONE: (215) 568-3100
TELEPHONE: (215) 568-3439
INFORMATION FOR SEQ ID NO: 89:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             282 GGAAGCAGCAGCAATGTCTG 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20 GGAAGCAGCAGTGACAG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; LENGTH: 20
; TYPE: NUCLEIC ACID
; STRANDEDNESS: SINGLE
; TOPOLOGY: LINEAR
; ANTI-SENSE: NO
US-07-977-284A-89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 85.0
Matches 17; Conservative
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                                                                                                                                                                      CITY: Ph
STATE: PA
COUNTRY:
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Sequence 33, Application US/09470443

Patent No. 6441156

GENERAL INFORMATION:

APPLICANT: Lerman, Michael I.

APPLICANT: Minna, John D.

APPLICANT: Minna, John D.

APPLICANT: Sekido, Yoshitaka

APPLICANT: Sekido, Yoshitaka

APPLICANT: Sekido, Yoshitaka

APPLICANT: Ming-Hui

APPLICANT: Ming-Hui

APPLICANT: Ming-Hui

APPLICANT: MI-OF INVENTION: Calcium Channel Compositions and Methods of Use Thereof

FILE REPERENCE: NIH-OS043

CURRENT APPLICATION NUMBER: US/09/470,443

CURRENT FILING DATE: 1999-12-22

EARLIER PILING DATE: 1998-12-32

NUMBER OF SEQ ID NOS: 114

SOSTWARE: PATENTIN Ver. 2.0
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APPLICANT: Nicholas M. Dean
APPLICANT: Susan F. Murray
TITLE CANT: Susan F. Murray
TITLE OF INVENTION: ANTISENSE MODULATION OF TRANSFORMING GROWTH FACTOR BETA
FILE REFERENCE: ISPH-0498
CURRENT APPLICATION NUMBER: US/09/661,753
CURRENT PILING DATE: 2000-09-14
EARLIER APPLICATION NUMBER: 60/154,546
EARLIER FILING DATE: 1999-09-17
NUMBER OF SEQ ID NOS: 68
SEQ ID NO 55
LENGTH: 20
                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                  DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.1%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 71; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                               Indels
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; OTHER INFORMATION: Antisense Oligonucleotide
US-09-661-753-55
                                                                                                                                                                                                                           Query Match
1.1%; Score 15.2; D
Best Local Similarity 85.0%; Pred. No. 71;
Matches 17; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                    282 GGAAGCAGCAGCAATGTCTG 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-661-753-55; Reguence 55, Application US/09661753; Patent No. 6436909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           621 CAGGGACCAGCTCCAGGAGC 640
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                                                                                                                                                                                                                                                                                                                                                               20 GGAAGCAGCAGCAGTGACAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence
TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 89
SEQUENCE CHARACTERISTICS:
LENGTH: 20
TYPE: NUCLEIC ACID
STRANDEDNESS: SINGLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 85.09
Matches 17; Conservative
                                                                                                                                   TOPOLOGY: LINEAR ANTI-SENSE: NO
                                                                                                                                                                              US-08-256-426B-89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 28
US-09-470-443-33
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LENGTH: 20
TYPE: DNA
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KENULT 29
US-09-470-443-43

Sequence 43, Application US/09470443

Sequence 43, Application US/09470443

Sequence 43, Application US/09470443

GENERAL INFORMATION:
APPLICANT: Lerman, Michael I.
APPLICANT: Minna, John D.
APPLICANT: Mei, Ming-Hui
APPLICANT: Sekida
APPLICANT: Sekida
APPLICANT: Sekida
APPLICANT: Ouh, Fuh-Mei
TITLE OF INVENITON: Calcium Channel Compositions and Methods of Use Thereof
FILE REFERENCE: NIH-05043
CURRENT APPLICATION NUMBER: US/09/470,443

CURRENT APPLICATION NUMBER: 60/114,359
EARLIER PILING DATE: 1999-12-22

EARLIER PILING DATE: 1999-12-30

NUMBER OF SEQ ID NOS: 114

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 43

LENGTH: 20

THENTH: 20

THENTH: 20
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                                                                                                                                                                                   Gaps
) ORGANISM: Artificial Sequence
; FEATURE:
) CTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-470-443-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
; CTHER INFORMATION: Description of Artificial Sequence: Synthetic US-09-470-443-43
                                                                                                                                                                                   0;
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Patent No. 6492170
GENERAL INFORMATION:
APPLICANT: Hong Zhang
APPLICANT: Andrew T. Watt
FILE REFERENCE: RTS-0183
CURRENT APPLICATION HOWBER: US/09/659,845A
CURRENT APPLICATION HOWBER: US/09/659,845A
CURRENT FILING DATE: 2001-07-23
SEQ ID NO: 174
SEQ ID NO: 168
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 1.1%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 71; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                DB 1; Length 20;
                                                                                                                                                                                   Indels
                                                                                                                             Score 15.2; DB Pred. No. 71; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Antisense Oligonucleotide US-09-659-845A-168
                                                                                                                                                                                                                            335 CIGGIGATAGICACAGIGGC 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                335 CTGGTGATAGTCACAGTGGC 354
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                                                                                                                             Query Match
Best Local Similarity 85.0%;
Matches 17; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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Gaps
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APPLICANT: Burton, Dennis R.
APPLICANT: Barbas, III, Carlos F.
APPLICANT: Chancek, Robert M.
APPLICANT: Murphy, Brian R.
APPLICANT: Murphy, Brian R.
APPLICANT: True, Jr., James E.
TITLE OF INVENTION: TO RESPIRATORY SYNCYTIAL VIRUS
                         CORRESPONDENCE ADDRESS:
ADDRESSEE: The Scripps Research Institute, Office of
ADDRESSEE: Patent Counsel
STREET: 10666 No. 5652138th Torrey Pines Road, Suite 220,
STREET: Mail Drop TPC8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Indels
                                                                                                                                                                                                                                                                                                      SOFTWARE PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/276,852
FILING DATE: 18-JUL-1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/178,302
FILING DATE: 30-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/954,148
FILING DATE: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R: Fish & Richardson P.C.
4225 Executive Square, Suite 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.2; D
Pred. No. 80;
0; Mismatches
                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 30, Application US/08162102C
; Patent No. 5762905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        262 CTGGGCTGGCTGATCAAAGA 281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21 credecrecerearchage 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-554-2937
TELEPAX: 619-554-6312
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                34,163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Ri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 92037
COMPUTER READABLE FORM:
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      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: La Jolla
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2
                                                                                                                                                                             USA
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US-08-162-102C-30/c
                                                                                                                                                        STATE: CA
COUNTRY: US
ZIP: 92037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ANTI-SENSE:
US-08-276-852-49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
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                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: GRIffais, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preve
TITLE OF INVENTION: and treatment of infection
FILE REPERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 655224

'Bequence 3952, Application US/09198452A

'Bequence 3952, Application US/09198452A

'Better No. 6552294

'GENERAL INFORMATION:

'TITLE OF INVENTION: Thereof and uses thereof, in particular for the diagnosis, prevering PILE REFERENCE: 9710-003-999

'CURRENT APPLICATION NUMBER: US/09/198,452A

'CURRENT APPLICATION NUMBER: US/09/198,452A

'CURRENT APPLICATION NUMBER: US/09/198,452A

'NUMBER OF SEQ ID NOS: 6849
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    Gaps
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Carlos F
Richard A
HUMAN NEUTRALIZING MONOCLONAL ANTIBODIES
TO HUMAN IMMUNODEFICIENCY VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
1.1%; Score 15.2; DB 1; Length 20;
Best Local Similarity 95.0%; Pred. No. 71;
Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.1%; Score 15.2; DB 1; Length 20;
BS.0%; Pred. No. 71;
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  0; Mismatches
                                                                                                                                                                    US-09-198-452A-1582/c
; Sequence 1582, Application US/09198452A
; Patent No. 6559294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 49, Application US/08276852
Patent No. 5652138
GENERAL INFORMATION:
APPLICANT: Burton, Dennis R
APPLICANT: Barbas, Carlos F
APPLICANT: Lerner, Richard A
TITLE OF INVENTION: HUMAN UBURALIZ
TITLE OF INVENTION: TO HUMAN IMMUNC
                                         567 ACTGCTCCAGCCAGGCCCTCC 586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       274 ATCAAAGAGGAAGCAGCAGC 293
                                                                                  1 ACTGCTCCAGAATGCCATCC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17 TGGATTAAACCAAACCCAGC 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 Archargcgaagcagc 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 rédarraraccadecadec 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: DNA
; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-3952
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-1582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17; Conservative
  Conservative
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US-08-276-852-49/c
  17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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Matches
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APPLICANT: Burton, Dennis R
APPLICANT: Burton, Dennis R
APPLICANT: Burton, Carlos F
APPLICANT: Lerner, Richard A
TITLE OF INVENTION: HUMAN NEUTRALIZING MONOCLONAL ANTIBODIES
TITLE OF INVENTION: TO HUMAN IMMUNODEFICIENCY VIRUS
NUMBER OF SEQUENCES: 170
CORRESPONDENCE ADDRESS:
ADDRESSEE: The Scripps Research Institute, Office of
ADDRESSEE: Patent Counsel
STREET: 10666 No. 5804440th Torrey Pines Road, Suite 220,
STREET: Mail Drop TPC8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
1.1%; Score 15.2; DB 1; Length 21;
Best Local Similarity 85.0%; Pred. No. 80;
Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER REDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: TBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
SOFTWARE: PATENTIN DATA:
APPLICATION NUMBER: US/08/899,575
FILLING DATE: 24-JUL-1997
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/276,852
FILING DATE: 18-UUL-1994
APPLICATION NUMBER: US 08/178,302
FILING DATE: 30-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US 07/954,148
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 49, Application US/08899575
Patent No. 5804440
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           262 CTGGGCTGGCTGATCAAAGA 281
                              FILING DATE: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: SCR14
TELEPHONE: 619-554-2937
TELEPHONE: 619-554-6112
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: mucleic acid
STRANDENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21 CTGGGCTGCCTGGTCAACGA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         34,163
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TELECOMMUNICATION INFORMATION
TELEPHONE: 619-554-2937
TELEFAX: 619-554-6312
                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
FILING DATE: 30-SEP-1992
ATTONEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,16
                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                        HYPOTHETICAL: N
HYPOTHETICAL: N
HAPOTHETICAL: NO
HAPOTHETICAL: NO
HAPOTHETICAL: NO
US-08-899-575-49
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Patent No. 5770440

GENERAL INFORMATION:
GENERAL INFORMATION:
TAPPLICANT: Barbas, Carlos F
APPLICANT: Barbas, Carlos F
APPLICANT: Lerner, Richard A
TITLE OF INVENTION: HUMAN NEUTRALIZING MONOCLONAL ANTIBODIES
TITLE OF INVENTION: 170 HUMAN IMMUNOBEFICIENCY VIRUS
NUMBER OF SEQUENCES: 170
CORRESPONDENCE ADDRESS:
ADDRESSEE: Patent Counsel
STREET: 10666 No. 5770440th Torrey Pines Road, Suite 220,
STREET: Mail Drop TPC8
CITY: La Jolla
STATE: CA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER P. 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Indels
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/162,102C
RILNG DATE: U-DEC-1993
CLASSIFICATION: 424
ATTONENTY/AGENT INFORMATION:
NAME: Halle, Ph.D., Lisa A.
REGISTRATION NUMBER: 38,347
REGISTRATION NUMBER: 38,347
REGISTRATION NUMBER: 38,347
REGISTRATION NUMBER: 30,347
RELEEVANCE (619) 679-5099
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
LYPE: NUCLeic acid
STRANDEDNES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/899,575
FLING DATE: 24-UU-1997
CLASSIFICATION 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/276,852
FILING DATE: 16-JUL-1994
APPLICATION NUMBER: US 08/178,302
FILING DATE: 30-SEP-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.1%; Score 15.2; I 85.0%; Pred. No. 80; Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           262 CTGGGCTGGCTGATCAAGA 281
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: DI
IMMEDIATE SOURCE:
CLONE: GD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 35
US-08-899-575-49/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    , LOCATION:
US-08-162-102C-30
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RESULT 38
US-08-635-109-21/c
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                                                                                                                                                                                                       DB 1; Length 21;
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                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Akitaya, Tatsuo
APPLICANT: Mitsuhashi, Masato
APPLICANT: Cooper, Alian
TITLE OF INVENTION: METHOD AND REAGENT
TITLE OF INVENTION: FOR MEASURING MESSENGER RNA
NUMBER OF SEQUENCES: 457
CORRESPONDENCE ADDRESS:
ADDRESSEE: Knobbe, Martens, Olson, and Bear
STREET: 620 Newport Center Dr. Sixteenth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SCTURENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/974,409C
FILING DATE: 12-NOV-1992
CLASSIFICATION: 435
ATTORNEY/AGENT INPORMATION:
NAME: Altman, Daniel B.
REGISTRATION NUMBER: 34,115
REBERRICE/POCKET UNBER: HITACHI.006CP2
TELEBCOMMUNICATION INPORMATION:
TELEPHONE: 714-760-0404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                     1.1%; Score 15.2; I
85.0%; Pred. No. 80;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                      US-07-974-409C-137/c
; Sequence 137, Application US/07974409C
; Patent No. 6300058
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 92660
COMPUTER READMBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   492 CGGTGTGCAGCGTCTTGGGG 511
                                                                                                                                                                                                                                                                                262 CTGGGCTGGCTGATCAAAGA 281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 ceretrcaecricireaee 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   137:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: CDNA to mRNA HYPOTHETICAL: NO
                                                                                                             DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 714-760-9502
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 85.0 nes 17; Conservative
               SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                           Conservative
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: doub
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                                                                                            linear
                                                                                                                                                                                                                     Best Local Similarity Matches 17, Conserv
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                                                                                        TOPOLOGY: lin
MOLECULE TYPE:
HYPOTHETICAL: N
                                                                                                                                                ; ANTI-SENSE: NO
US-08-899-575-49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE:
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                                                                                                                                                                                                       Query Match
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Sequence 137, Application PC/TUS9300977

GENERAL INFORMATION: METHOD AND REAGENT FOR MEASURING MESSENGER RNA TITLE OF INVENTION: METHOD AND REAGENT FOR MEASURING MESSENGER RNA NUMBER OF SEQUENCES: 711

CORRESPONDENCE ADDRESS:
ADDRESSEE: Knobbe, Martens, Olson, and Bear STREET: 620 Newport Center Dr. Sixteenth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .
0
                                        GENERAL INFORMATION:
APPLICANT: Persson, Mats A. A.
APPLICANT: Parsson, Mats E.
APPLICANT: Allander, Tobias E.
TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES SPECIFIC FOR
TITLE OF INVENTION: HEPATITIS C VIRUS (HCV) E2 ANTIGEN
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: REED & ROBINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 21;
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SOFTWARE: Patentin Release #1.0, Version #1.25
SURINARE: APPLICATION DATA:
APPLICATION UNMERS: PCT/US93/00977
FILING DATE: 19930129
                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
RPLING DATE: 19-APR-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.1%; Score 15.2; 1
85.0%; Pred. No. 80;
                                                                                                                                                                                                      3: REED & ROBINS
285 Hamilton Avenue, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: MCCACKEN, Thomas P
REGISTRATION NUMBER: 38,548
REGISTRATION NUMBER: 330-6146
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 327-3400
TELEFRA: (415) 327-3231
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            262 CTGGGCTGGCTGATCAAAGA 281
; Sequence 21, Application US/08635109
; Patent No. 6538114
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US-08-635-109-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 19-APR-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 85.09
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                         STREET: 285 Hamilt
CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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Gaps

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APPLICANT: Zhang, Lin
APPLICANT: Zhang, Wei
TITLE OF INVENTION: Gene Expression Profiles in No. 6333152mal and
TITLE OF INVENTION: Gancer Cells
FILE REFERENCE: 01107.74664
CURRENT APPLICATION NUMBER: US/09/081,646
CURRENT FILING DATE: 1998-05-20
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ ID NOS: 871
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 727
LENGTH: 15
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Sequence 2539, Application US/08585684B

Patent No. 5877021

GENERAL INFORMATION:

APPLICANT: Stinchcomb, Daniel T.

APPLICANT: MCSWiggen, James

TITLE OF INVENTION: METHOD AND REAGENT FOR THE

TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE

TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES

NUMBER OF SEQUENCES: 2751

CORRESPONDENCE ADDRESS:

ADDRESSEE: Lyon & Lyon

STREET: 631 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 15; 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.5
COMPUTER: HER COMPATION DATA:
APPLICATION NUMBER: US/08/585,684B
APPLICATION NUMBER: US/08/585,684B
                                                                                                                                                                                                                                                                                                                                                                                                        Score 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: January 16, 1996,
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/000,951
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
                                                                                                                                                                                                                                                                                                                                                                                     1.1%; Scc.
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REPRENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 745 CAIGITGCIGACITI 759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 1.1
Best Local Similarity 100.
Matches 15; Conservative
  APPLICANT: Vogelstein, Bert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 18 base pairs
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                            TYPE: DNA
CORGANISM: Homo sapiens
US-09-081-646-727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
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TO HUMAN IMMUNODEFICIENCY VIRUS
170
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                                                                                                                                                                                                                                                                                                                                                                                                                                Length 21;
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AFFLICANT;
TITLE OF INVENTION: HUMAN NEUTRALIZING MONOCLONAL ANTIBODI
TITLE OF INVENTION: TO HUMAN IMMUNOBERICIENCY VIRUS
NUMBER OF SEQUENCES: 170
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
FILING DATE: 11-JUL-1995
PRIOR APPLICATION DATA:
FILING DATE: 18-JUL-1994
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 1.1%; Score 15.2; Dest Local Similarity 85.0%; Pred. No. 80; Best Local 77; Conservative 0; Mismatches
                                          NAME: Altman, Daniel E.
REGISTRATION NUMBER: 34,115
REFERENCE/DOCKET NUMBER: HITACHI.006H
TELECOMMUNICATION INFORMATION:
TELEPHONE: 714-760-0404
TELEFAX: 714-760-9502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 40
PCT-US95-08743-49/C
Sequence 49, Application PC/TUS9508743
Seguental INFORMATION:
APPLICANT:
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US-09-081-646-727
Sequence 727, Application US/09081646
Patent No. 6333152
GENERAL INFORMATION:
APPLICANT: Kinzler, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            262 CTGGGCTGGCTGATCAAAGA 281
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                                                                                                                                                                                                                                                                                                                    CDNA to mRNA
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Altman, Daniel E.
                                                                                                                                                    TELEFAX: 714-760-9502
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                        LENGTH: 21
TYPE: NUCLEIC ACID
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                    8
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MOLECULE TYPE: 1
HYPOTHETICAL: N
                                                                                                                                                                                                                                                                                                                                                         ; ANTI-SENSE: NO PCT-US93-00977-137
                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCT-US95-08743-49
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RESULT 45
US-08-714-991-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Patent No.
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                                                                  DB 1; Length 18
                                                                                                                                                                                                                                                                                                                                                                             RESULT 43
US-09-038-073-2539/C
US-09-038-073-2539/C
Sequence 2539, Application US/09038073
Seque
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                                                                                                                                        2; Indels
                                                                  1.1%; Score 14.8; D
88.9%; Pred. No. 67;
iive 0; Mismatches
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88.9%; Pred. No. 67;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FASTEEQ Version 1.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/038,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   218/078
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US-08-630-592-14
Sequence 14, Application US/08630592
Patent No. 5770432
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/585,684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REEBENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 989-1600
TELERS: (5-3510
INFORMATION FOR SEQ ID NO: 2539:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: mucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      277 AAAGAGGAAGCAGCA 294
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                                                                  Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
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Best Local Similarity
Matches 16; Conserv
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US-08-585-684B-2539
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 90071
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STREET:
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Gaps
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APPLICANT: NORTH, Michael
APPLICANT: NISHINA, Patsy
APPLICANT: NO. 5776762en-Trauth, Konrad
APPLICANT: NAGGERT, Juergen
TITLE OF INVENTION: OBESITY ASSOCIATED GENES
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: 4 Embarcadero Center, Suite 3400
APPLICANT: Nishina, Patsy
APPLICANT: No. 5770432enTrauth, Konrad
APPLICANT: Naggert, Juergen
APPLICANT: No. 5770432th, Michael
TITLE OF INVENTION: Obesity Associated Genes
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS: ADDRESSE: FLERR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: 3400 Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PCDCS/MSDOS
SOCTWARE: Patentl N. Release #1.0, Version #1.30
CURRENT APELICATION DATA:
APPLICATION NUMBER: US/08/630,592
FILING DATE:
FILING DATE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J.
REGISTRATION NUMBER: 36,677
REFERENCE/DOCKET NUMBER: 35,677
TELEFENCE/DOCKET NUMBER: 35,677
TELEFENCE/DOCKET NUMBER: 35,677
TELEFENCE/DOCKET NUMBER: 35,677
TELEFENCE/DOCKET NUMBER: 35,677
TELEFENCE/MONISCRIPTION INFORMATION:
TELEFENCE (415) 7811989
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.8; Pred. No. 76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "primer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 14, Application US/08714991
Patent No. 5776762
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Best Local Similarity 88.9%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                             Floppy disk
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 4 Embarcadero
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                      ZIP: 941114187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                  STREET: 3400 Embarce
CITY: San Francisco
STATE: California
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822 CCTGATGCAGCTGAAGCT 839
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (213) 489-1600
                                                                                 Query Match
Best Local Similarity 88.3%
...hes 16; Conservative
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Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
           single
                         TOPOLOGY: linear MOLECULE TYPE: CDNA
         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE:
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US-08-623-891-3/c
                                                              US-09-032-365A-26
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Patent No. 6114502

GENERAL INFORMATION:
APPLICANT: No. 6114502th, Michael
APPLICANT: Nishina, Patsy
APPLICANT: Nagaart, Juergen
APPLICANT: No. 6114502en-Trauth, Konrad
TITLE OF INVENTION: GENE FAMILY ASSOCIATED WITH
TITLE OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
APPLICAL-
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: SHERWOOD, Pamela J.
REGISTRATION NUMBER: 36,677
; REFERENCE/DOCKET NUMBER: A-59504-1/PJS
; TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-494-8700
"FLEFAX: 415-494-8770
"FLEFAX: 415-494-8771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/032,365A
                                                                                                                                                                                                                                                                                                                                                                                               1.1%; Score 14.8; D
88.9%; Pred. No. 76;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Boziceric & Reed, LLP
STREET: 285 Hamilton Avenue, Suite 200
CIIY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,677
REFERENCE/DOCKET NUMBER: SEQ-2CIP2
TELEPHONE: 650-327-3400
                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "primer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         822 CCTGATGCAGCTGAAGCT 839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 CCTGAGGCAGCAGAAGCT 19
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Best Local Similarity 88.9
Matches 16; Conservative
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SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                             LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 650 327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94301
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       S
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US-09-032-365A-26
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                                                          Gaps
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1.1%; Score 14.8; DB 1; Length 19;
88.9%; Pred. No. 76;
Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                         Sequence 3, Application US/08623891
; Patent No. 579578
; GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: INHIBITING HERPES SIMPLEX:
TITLE OF INVENTION: INFORMATION
NUMBER OF SEQUENCES: 115
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREE: 611 West Sixth Street
CITY: Los Angeles
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: Wordberfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/623,891
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION: 435
PRIOR PAPILCATION DATA:
APPLICATION NUMBER: US/08/238,200
FILING DATE:
APPLICATION NUMBER: US/07/987,133
FILING DATE:
APPLICATION NUMBER: US/07/987,133
FILING DATE: May 14, 1992
APPLICATION NUMBER: 07/982,921
FILING DATE: September 18, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warbley, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 200/209
TELECOMINICATION INFORMATION:
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                       FILING DATE:
CLASSIFICATION:
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US-09-634-262-3/c
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                                                  Sequence 42, Application US/09286904A
| Patent No. 6140124
| Patent No. 6140124
| Patent No. 6140124
| APPLICANT: Monia, Brett P.
| APPLICANT: Monia, Brett P.
| APPLICANT: Moss, Pamela S.
| APPLICANT: McKay, Robert
| APPLICANT: McKay, Robert
| TITLE OF INVENTION: Activated Protein Kinase Expression
| TITLE OF INVENTION: Activated Protein Kinase Expression
| FILE REFERENCE: ISPH-0347
| CURRENT APPLICATION NUMBER: US/09/286,904A
| CURRENT PILING DATE: 1999-04-06
| NUMBER OF SEQ ID NOS: 95
| SOGTWARRE: Patentin Ver. 2.0
| SEQ ID NO 42
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APPLICANT: Patrick Allen Marcotte
APPLICANT: Lex M. Cowsert
TITLE OF INVENTENCE: RE-0090
CURRENT APPLICATION NUMBER: US/09/742,703
CURRENT FILING DATE: 2000-12-20
NUMBER OF SEQ ID NOS: 49
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
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APPLICANT: Kenneth G. Draper
ITILE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: INHIBITING HERRES SIMPLEX
TITLE OF INVENTION: VIRUS REPLICATION
NUMBER OF SEQUENCES: 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ) OTHER INFORMATION: Antisense Oligonucleotide US-09-742-703-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.1%; Score 14.8; D
88.9%; Pred. No. 87;
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
1.1%; Score 14.8; E
Best Local Similarity 88.9%; Pred. No. 87;
Matches 16; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ) OTHER INFORMATION: antisense sequence US-09-286-904-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 49
US-09-742-703-11
Sequence 11, Application US/09742703
; Patent No. 6423543
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 3, Application US/09340861
Patent No. 6432704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1287 TACAGTTGCTCAGCCTGG 1304
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                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                 RESULT 48
US-09-286-904-42/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 50
US-09-340-861-3/c
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Gaps
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1.1%; Score 14.8; DB 1; Length 20;
Best Local Similarity 88.9%; Pred. No. 87;
Matches 16; Conservative 0; Mismatches 2; Indels
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Patent No. 6440719;
GENERAL INFORMATION:
APPLICANT: Kenneth G. Draper TITLE OF INVENTION: INHIBITING HERPES SIMPLEX;
TITLE OF INVENTION: INHIBITING HERPES SIMPLEX;
TITLE OF INVENTION: VIROS REPLICATION
NUMBER OF SEQUENCES: 115
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: 1BM Compatible
OPERATING SYSTEM: 1BM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/340,861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRICE APPLICATION DATA:
APPLICATION NUMBER: US/07/987,133
FILING DATE:
APPLICATION NUMBER: US/07/987,133
FILING DATE: MAY 14, 1992
APPLICATION NUMBER: 07/948,359
FILING DATE: September 18, 1992
ATTORNEY,AGENT INFORMATION:
NAME: WATDEL WATDEL SEPTEMBER: 200/209
FREISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 200/209
TELECHANGING CONTROL OF TELECHANGING CONTROL CONTROL
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APPLICATION NUMBER: US/09/634,262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E: Lyon & Lyon
611 West Sixth Street
ADDRESSEE: Lyon & Lyon
STREET: 611 West Sixth Street
CITY: Los Angeles
STATE: California
COUWIRY: USA
ZIP: 90017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        130 GGACAGGGACGCCCGCTC 147
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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MEDIUM TYPE: 3.5" Di
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US-09-359-921-27/c
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TELEX: 6'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: McRay, Robert
APPLICANT: McRay, Robert
TITLE OF INVENTION: Antisense Modulation of p38 Mitogen
TITLE OF INVENTION: Antisense Modulation of p38 Mitogen
TITLE OF INVENTION: Activated Protein Kinase Expression
FILE REFERENCE: ISPH-0488
CURRENT APPLICATION NUMBER: US/09/640,101
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: 09/286,904
PRIOR FILING DATE: 1999-04-06
NUMBER OF SEQ ID NOS: 107
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
1.1%; Score 14.8; DB 1;
Best Local Similarity 88.9%; Pred. No. 87;
Matches 16; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                      APPLICATION NUMBER: 07/882,921
FILING DATE: May 14, 1992
APPLICATION NUMBER: 07/948,359
FILING DATE: September 18, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 20,327
REFERENCE/DOCKET NUMBER: 20,327
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: (213) 955-0440
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: antisense sequence
                     US/07/987,133
                                                                                                                                                                                                           200/209
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 42, Application US/09640101 Patent No. 6448079
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19 GGACAGGGACGGCCGATC
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STRANDEDNESS: single
TOPOLOGY: linear
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 52
US-09-640-101-42/c
                                                                                                                                                                                                                                                                                                                                                       LENGTH: 20
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US-09-099-053-19
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APPLICANT: DERVAN, PETER B.
APPLICANT: DERVAN, BAIRD, ELDON B.
TITLE OF INVENTION: METHOD FOR THE SYNTHESIS OF PYRROLE AND IMIDAZOLE TITLE OF INVENTION: CARBOXAMIDES ON A SOLID SUPPORT FILE REFERENCE: 025098-1602
CURRENT APPLICATION NUMBER: US/09/359,921
CURRENT FILING DATE: 1999-07-22
NUMBER OF SEQ ID NOS: 31
SOFTWARE: PATENTING DATE: 2.1
SEQ ID NO 27
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                           APPLICANT: Sara Courtheldge
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF
TITLE OF INVENTION: SAD RELATED DISORDERS
TUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEB: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8;
Pred. No. 98
                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/099,053
FILING DATE: Herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 27, Application US/09359921
Patent No. 6545162
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     60/049,914
                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       374 CCCAGCTTCCTCCAGAGG 391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327 REFERENCE/DOCKET NUMBER: 23 TELECOMMUNICATION INFORMATION: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/049,
FILING DATE: June 18, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 67.3510
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (213, 955-0440
Greg Plowman
Susan Onrust
David Markby
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16; Conservative
                                                                                                                                                                                                                                       CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
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; TYPE: DNA; ORGANISM: HUMAN US-09-177-776-113
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                                                                                    1.1%; Score 14.4; DB 1; Length 17; 93.8%; Pred. No. 71;
                                                                                                                                                    1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HARLER INO. 82.7041

APPLICANT: Zavada, Jan
APPLICANT: Zavada, Jan
APPLICANT: Zavada, Jan
APPLICANT: Pastorekova, Silvia
APPLICANT: Pastorekova, Silvia
APPLICANT: Pastorekova, Silvia
APPLICANT: Pastorekova, Jaromir
TITLE OF INVENTION: MN Gene and Protein
FILE REPERENCE: D-0021.5A
CURRENT FILING DATE: 1998-10-23
EARLIER APPLICATION NUMBER: 09/177, 776
EARLIER FILING DATE: 1998-00-24
EARLIER FILING DATE: 1995-06-07
EARLIER FILING DATE: 1995-06-
                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 113, Application US/09178115 ; Patent No. 6297041
OTHER INFORMATION: oligonucleotide
                                                                                                                                                 0;
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                                                                                                                                                 Conservative
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Best Local Similarity
"Ahas 15; Conserv
                                                                                                                 Best Local Similarity
Matches 15; Conserv
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US-09-177-776-113/c
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; ORGANISM: HUMAN
US-09-178-115-113
                                                                                                                                                                                                                                                                                                                                                                RESULT 55
US-09-178-115-113/c
      ; OTHER INFORM
US-09-359-921-27
                                                                                       Query Match
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APPLICANT: Li, Xiao-Jiang
APPLICANT: Blackshaw, Seth
APPLICANT: Blackshaw, Seth
APPLICANT: Bryder, Solomon H.
TITLE OF INVENTION: AMILORIDE-SENSITIVE SODIUM CHANNEL AND
TITLE OF INVENTION: SALTY TASTE PERCEPTION
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Allegretti, LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ó
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/376,362A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 1001 G Street, N.W., Eleventh Floor CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.1%; Score 14.4; I 93.8%; Pred. No. 82;
                                                    CURRENT APPLICATION NUMBER: US/09/177,776A
                                                                      CURRENT FILING DATE: 1998-10-23
EARLIER APPLICATION NUMBER: 09/787,739
EARLIER FILING DATE: 1998-10-23
EARLIER FILING DATE: 1995-06-07
EARLIER APPLICATION NUMBER: 08/485,863
EARLIER APPLICATION NUMBER: 08/487,077
EARLIER FILING DATE: 1995-06-07
EARLIER FILING DATE: 1995-06-15
EARLIER FILING DATE: 1995-06-15
EARLIER FILING DATE: 1995-06-15
EARLIER FILING DATE: 1995-06-15
EARLIER FILING DATE: 1992-10-21
EARLIER FILING DATE: 1992-10-21
EARLIER FILING DATE: 1992-10-21
EARLIER FILING DATE: 1992-03-11
NUMBER OF SEO ID NOS: 116
TITLE OF INVENTION: MN Gene and Protein FILE REFERENCE: D-0021.5A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-376-362A-8/c
; Sequence 8, Application US/08376362A; Patent No. 5693756
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A.
ZIP: 20001-4597
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk

COMPUTER: ISM PC COMPACTIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1017 GAGATGGTGCCAAAGT 1032
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Best Local Similarity 93.8
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM IIEE FLOURY MIEE.

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/08/450,905B
FILING DATE: 26-MAR-1995
PRIOR APPLICATION NUMBER: 07/982,759
PRIOR APPLICATION NUMBER: 07/982,759
PRIOR APPLICATION NUMBER: 07/982,759
FILING DATE: 23-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9127319.3
FILING DATE: 23-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9221587.0
FILING DATE: 14-OCT-1992
ATTOMENYAGEN: LINCOMMATION:
ANAME: MANDER LINCOMMATION:
                                                                                                                                                                                           Sequence 134, Application US/08450905B
Patent No. 5856301
GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: Stem Cell Inhibiting Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc_feature
LOCATION: 1..20
OTHER INFORMATION: /product= "BB9513 oligomer"
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93.8%; Pred. No. 1.1e+02;
tive 0; Mismatches 1;
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Patent No. 6057123
GENERAL INFORMATION:
APPLICANT: CRAIG, Stewart
APPLICANT: GEORGE, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 102...
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-526-6110
TELEFAX: 617-526-5000
INFORMATION FOR SEQ ID NO: 134:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31,321
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                                             670 TTGGCCAGCGTGTAT 685
                                                                                         3 TAGGCCAGCGTGGTAT 18
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                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 178
CORRESPONDENCE ADDRESS:
ADDRESSES: HALE and DORR
STREET: 60 State Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: BAKER, HOLLIE L. REGISTRATION NUMBER: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 93.8
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-450-905B-134
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                                                                                                                                                         RESULT 59
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                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 1.1%; Score 14.4; DB 1; Length 20; Best Local Similarity 93.8%; Pred. No. 1.1e+02; Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
CIVIRX: USA
ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: HALVERSON, Joy L.
APPLICANT: DAVORAK, Jan
TITLE OF INVENTION: AVIAN SEX IDENTIFICATION PROBES
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/634,331
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FILING DATE: 23-UAR - CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Kagan A., Sarah
REGISTRATION NUMBER: 32,141
REFRENCE/DOCKET NUMBER: 01107
TELECOMMUNICATION INFORMATION:
TELEFAX: 202-508-929
INFORMATION FOR SEQ ID NO: 8:
IENGTH: 20 base pairs
ILENGTH: 20 base pairs
ILENGTH: 20 base pairs
TYPE: mucleic acid
STRANDENNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
US-08-376-362A-8
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US-08-634-331-3
Sequence 3, Application US/08634331
; Patent No. 5707809
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TELEFAX: (415) 494-8771
TELEX: 910 2777299FHT UR
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: mucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         629 AGCTCCAGGAGCTCTG 644
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Best Local Similarity 93.8
Matches 15; Conservative
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DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-634-331-3
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Lee, Cheng-Chi
APPLICANT: Albrecht, Urs
APPLICANT: Albrecht, Urs
APPLICANT: Bichele, Gregor
APPLICANT: Bichele, Gregor
FILE APPLICANT: Bichele, Gregor
TITLE OF INVENTION WAMMAIIAN CIrcadian Rhythym-Like Gene
CURRENT APPLICATION NUMBER: US/09/150,460B
CURRENT APPLICATION NUMBER: US 60/058,256
FRIOR FILING DATE: 1998-09-09
FRIOR FILING DATE: 1997-09-09
NUMBER OF SEQ ID NOS: 21
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
1.1%; Score 14.4; DB 1; Length 20;
Best Local Similarity 93.8%; Pred. No. 1.1e+02;
Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
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Best Local Similarity 93.8%; Pred. No. 1.1e+02;
Matches 15; Conservative 0; Mismatches 1;
                                                                      SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Lidata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-034
TELEPHONE: 609-810-1155
TELEPHONE: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 2, Application US/09150460B; Patent No. 6190882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 7, Application US/09228942; Patent No. 6203988; GENERAL INFORMATION:
                                                  WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        624 GGACCAGCTCCAGGAG 639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            788 CCAGTGCCCTGGCTCG 803
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-48
                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 63
US-09-228-942-7/c
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US-09-150-460B-2
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Fatent No. 6184212
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Batentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/982,759F
FILING DATE: 23-DEC-1991
PRIOR APPLICATION NUMBER: GB 9221587.0
FILING DATE: 14-OCT-1992
ATTORNEY/AGENT INFORMATION:
REPRESENCE/DOCKET NUMBER: 102378.120
FILING DATE: 31.321
REPRESENCE/DOCKET NUMBER: 102378.120
TELEROMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEROMMUNICATION INFOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
APPLICANT: EDWARDS, Richard Mark
APPLICANT: CZAPLEWSKI, Lloyd George
APPLICANT: GILBERT, Richard
TITLE OF INVENTION: Stem Cell Inhibiting Proteins
NUMBER OF SEQUENCES: 178
CORRESPONDENCE ADDRESS:
ADDRESSEE: HALE and DORR LLP
STREET: 60 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc feature
LOCATION: 1..20
OTHER INFORMATION: /product= "BB9513 oligomer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                840 TTCAGATGGGTCAGCA 855
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                            ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U.S.A.
                                                                                                                                                                                                 ADDALL
STREET: 60
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STATE: MA
7. 02109
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US-09-280-805-48/c
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ZIP: 08053
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Query Match
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                                                                                                                                                                                                                                                   FEATURE:
O'THER INFORMATION: Description of Artificial Sequence:
CHER INFORMATION: oligonucleotide ligated to 3' end of DNA fragment
US-09-228-942-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 7, Application US/08246489
; Fatent No. 6225049
; GENERAL INFORMATION:
APPLICANT: Lan, Michael S.
APPLICANT: No. 6225049kins, Abner L.
TITLE OF INVENTION: NOVEL HUMAN INSULINOMA-ASSOCIATED CDNA
NUMBER OF SEQUENCES: 27
NUMBER OF SEQUENCES:
ADDRESSEE: Knobbe, Martens, Olson & Bear
STREET: 620 Newport Center Drive
CITY: Newport Beach
STATE: California
COUNTRY: USA
ZIP: 9266
APPLICANT: Unmartat, Allihiro
TITLE OF INVENTION: DNA FRAGMENT ANALYSIS METHOD AND REAGENT KIT
TITLE OF INVENTION: DNA FRAGMENT ANALYSIS METHOD AND REAGENT KIT
CURRENT APPLICATION UNMBER: US/09/228,942
CURRENT FILING DATE: 1999-01-12
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 7
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 240, Application US/09517467B
Fatent No. 6451602
GENERAL INFORMATION:
APPLICANT: Lax M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF PARP EXPRESSION
FILE REFERENCE: RIS. 0150
CURRENT APPLICATION NUMBER: US/09/517,467B
CURRENT APPLICATION NUMBER: 09/517,467B
FRIOR APPLICATION NUMBER: 09/517,467
FRIOR APPLICATION NUMBER: 09/517,467
FRIOR FILING DATE: 2000-03-02
FRIOR FILING DATE: 345
                                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.1%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.1e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                              Query Match
1.1%; Score 14.4; DB 1;
Best Local Similarity 93.8%; Pred. No. 1.1e+02;
Matches 15; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                         1144 ITTTTTTTTTGGA 1159
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                                                                                                                                                                                                        TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18 Triririririgga 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15; Conservative
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 64
US-09-517-467B-240
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US-08-246-489-7
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LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18, Application US/08033081B
Fatent No. 549851
GENERAL INFORMATION:
TITLE OF INVENTION: DISCREDITARY RETINAL
TITLE OF INVENTION: DISCREDENCE:
TITLE OF INVENTION: DISCREDENCE:
TITLE OF INVENTION: DESCRIBATIVE DISCRESSONERCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: ADDRESSE:
ADDRESSEE: ADDRESSE:
ADDRESSEE: ADDRESSE:
ADDRESSEE: ABONENCE ADDRESSE:
ATTE: MASSACHUSELTS
COUNTRY: U.S.A.
TITLE OF INVENTION STATES
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM PS/2 MOdel 50Z or 55XX
OPERATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARR: WORDERFECT (Version 5.1)
                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/246,489
                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US/07/901,715
FILING DATE:
ATORNEY/AGENT INFORMATION:
MAME: Israelsen, Ned A.
REGISTRATION NUMBER: 29,655
REFERENCE/DOCKET NUMBER: 19,655
REPERENCE/DOCKET NUMBER: NIH012.012A
TELECOMMUNICATION INFORMATION:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/033,081B
FILING DATE: March 11, 1993
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: January 23, 1992
APPLICATION NUMBER: 07/469,215
FILING DATE: January 24, 1990
ATTORNEY, AGENT INFORMATION:
NAME: CLARK, PRUL T.
REGISTRATION NUMBER: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07/825,296
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 84.2%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: March 11,
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07
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STRANDEDNESS: single
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                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-033-081B-18
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APPLICANT: D'OVIDIO, RENATO
APPLICANT: POCCEDOU, ENRICO
APPLICANT: POCCEDOU, ENRICO
APPLICANT: PRECHITELLI, CINZIA
APPLICANT: CARDELLI, LUISA ERCOLI
TITLE OF INVENTION: ISOLATION AND CHARACTERIZATION OF A GENE
TITLE OF INVENTION: ENCODING A LOW MOLECULAR WEIGHT GLUTENIN
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
STREET: 1755 S. JEFFERSON DAVIS HIGHMAY
CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ô
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.0%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 1.2e+02; Live 0; Mismatches 3; Indels
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MEDIUM TYPE: Diskette
COMPUTER: DISKette
COMPUTER: DISKette
COMPUTER: DISKETE
COMPUTER: DISKET
COMPUTER: DIS
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Patent No. 5952202
GENERAL INFORMATION:
            1159 AAGTAAAGCAGCTAAAACA 1177
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; Patent No. 5973225
; GENERAL INFORMATION:
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TELEFAX: (650) 638-671
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 DARGET CONTRACTOR CONTRACT
                                                                                                        1 AAGTAAAGCCGCAAAAGCA 19
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Best Local Similarity 84.2'
Matches 16; Conservative
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STRANDEDNESS: single
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| Patent No. 581760
| GENERAL INFORMATION:
| APPLICANT: Evans, Glen A. |
| APPLICANT: Smitch, Michael W. |
| TILLE OF INVENTION: METHOD FOR GENERATION OF SEQUENCE |
| TILLE OF INVENTION: METHOD FOR GENERATION OF SEQUENCE |
| TILLE OF INVENTION: METHOD FOR GENERATION OF SEQUENCE |
| TILLE OF INVENTION: METHOD FOR GENERATION OF SEQUENCE |
| TILLE OF INVENTION: METHOD MAPS OF COMPLEX GENOMES |
| CORRESPONDENCE ADDRESS: | ADDRESSEE: Pretty, Schroeder, Brueggemann & Clark |
| STREET: Los Angeles |
| STATE: CA AN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CAPUTER FALABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Ralease #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/117,952
FILING DATE: 07-SEP-1993
CLASSIFICATION NUMBER: US/08/117,952
FILING DATE: 15-JUN-1993
ATTORNEY/AGENT INFORMATION:
NAME: RELEATION NUMBER: 31,192
REGISTRATION STEPPINGER: 31,192
REGISTRATION STEPPINGER: 619-546-4737
TELEFRAM: 619-546-4737
INFORMATION FOR SEQ ID NO: 417:
SEQUENCE CHARACTERISTICS:
LUMBGTH: 20 DASE PAIRE
      00246/069005
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                                                 TELECOMMUNICATION INFORMATION
TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-8906
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                             TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single
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STRANDEDNESS: sing
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US-08-033-081B-18
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HYPOTHETICAL: NO
HATI-SENSE: NO
US-08-117-952-417
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US-08-117-952-417
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GENERAL INFORMATION:

APPLICANT: Futreal, Phillip A

APPLICANT: Mooster, Richard F

APPLICANT: Mooster, Alan

APPLICANT: Stratton, Michael R

TITLE OF INVENTION: Materials and methods relating to the

TITLE OF INVENTION: identification and sequencing of the BRCA2 cancer

TITLE OF INVENTION: susceptibility gene and uses thereof.

NUMBER OF SEQUENCES: 222

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Bell Seltzer Park & Gibson
STREET: 310 UCB Plaza, 3605 Glenwood Avenue, PO Drawer 31107
                                                                                                                                                                                                                                                               Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.0%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 1.2e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Petentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/755,587
FILING DATE: 25-NOV-1996
RIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9523959.6
FILING DATE: 23-NOV-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 952555.0
FILING DATE: 14-DEC-1995
FILING DATE: 14-DEC-1995
FILING DATE: 28-AUG-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                        1.0%; Score 14.2; DB 1; Length 2
84.2%; Pred. No. 1.2e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                         MOLECULE TYPE: other nucleic acid HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                      1144 TTTTTTTTTTGGAAGT 1162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 59, Application US/08755587 Patent No. 6045997
                                                                                                                                                                                                                                                                                                                                                                                                    TTTTTTTTTTTTTCCAGGT 20
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 31, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kenneth D Sibley
  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                      16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
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1.0
Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  : 20 base pairs
nucleic acid
                                                  20 base pairs
                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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                                                                                                                                                                                       ; ANTI-SENSE: NO
US-08-715-461-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-755-587-59/c
                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-755-587-59
                                                  LENGIH:
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Patent No. 598556
GENERAL INFORMATION:
APPLICANT: KAMBARA, Hideki
APPLICANT: OKANO, Kazunori
TITLE OF INVENTION: DNA SEQUENCING METHOD AND DNA SAMPLE
TITLE OF INVENTION: PREPARATION METHOD
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: ANTOMELLI, TERRY STOUT & KRAUS
STATE: VA
STATE: VA
STATE: VA
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
SOFTWARE: PatentIn Release #1.0, Version #1.30
SOFTWARE: PatentIn Release #1.0, Version #1.30
SOFTWARE: APPLICATION DATA:
APPLICATION NUMBER: US/08/715,461
FILING DATE: 18-SEP-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: TERRY, David T.
REGISTRATION NUMBER: 20,178
REGISTRATION NUMBER: 500.34872X00
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFHONE: 703-312-6600
                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: PatentIn Pelease #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/991,300
FILING DATE: 16-DEC-1997
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
TRING DATE: 19-DEC-1996
ATTORNEY/AGENT INFORMATION:
NAME: OBLOW, NORMAN F:
REGISTRATION NUMBER: 24,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
1.0%; Score 14.2; DB 1;
Best Local Similarity 84.2%; Pred. No. 1.2e+02;
Matches 16; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                               2264-0201-0X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "PRIMER"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1011 GCACCTGAGATGGTGCCAA 1029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 GCACCGGAGTTGGTGCCTA 2
                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
TELEFAX: 703-413-2220
                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
                        22202
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COUNTRY:
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US-08-715-461-4
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US-09-288-461-68/c
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                                                                                                                                                                                            APPLICANT: MCKAY, Robert A.
APPLICANT: MCKAY, Robert A.
APPLICANT: Dean, Nicholas M.
APPLICANT: Dean, Nicholas M.
APPLICANT: Dean, Nicholas M.
APPLICANT: Brett
APPLICANT: Brett
APPLICANT: Oande, William A.
TITLE OF INVENTION: POR THE WODULATION OF JNK PROTEINS
TITLE OF INVENTION: POR THE WODULATION OF JNK PROTEINS
FILE REPERENCE: 1SPH-0350
CURRENT APPLICATION NUMBER: US/09/287,796A
CURRENT PILING DATE: 1999-04-07
BARLIER PILING DATE: 1999-04-07
BARLIER PILING DATE: 1999-08-07
BARLIER PILING DATE: 1999-08-03
NUMBER OF SEQ ID NOS: 165
SEQ ID NO 123
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 16, Application US/09288461
Patent No. 6159694
GENERAL INFORMATION:
APPLICANT: Karras, James G.
TITLE OF INVENTION: Attisense Oligonucleotide Modulation of STAT3
TITLE OF INVENTION: Expression
FILLE REFERENCE: 18PH-0338
CURRENT APPLICATION NUMBER: US/09/288,461
CURRENT FILING DATE: 1999-04-08
NUMBER OF SEQ ID NOS: 107
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 16
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Synthetic Sequence US-09-287-796-123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Synthetic Sequence
                                                                                                                                         Sequence 123, Application US/09287796A Patent No. 6133246 GENERAL INFORMATION:
1110 AGTTTTCTGTTTAATTGAA 1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              628 CAGCICCAGGAGCICIGCA 646
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                                  20 AGICCICITITAGIIGAA 2
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 84.2%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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Matches 16; Conserv
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                                                                                                RESULT 72
US-09-287-796-123
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GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Madeline M. Butler
APPLICANT: Jacqueline Wyatt
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF PEPCK-CYTOSOLIC EXPRESSION
FILE REFERENCE: RTS-0123
CURRENT APPLICATION NUMBER: US/09/488,671A
NUMBER OF SEQ ID NOS: 177
SEQ ID NO 52
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 123, Application US/09130616C
Sequence 123, Application US/09130616C
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Moria, Sectt
APPLICANT: Moria, Brett
APPLICANT: Moria, Brett
APPLICANT: Gaarde, William A.
APPLICANT: Garde, William A.
APPLICANT: Garde, William A.
APPLICANT: Garde, William A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS AND METHODS
FILE REPERENCE: ISPH-0319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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Squence 68, Application US/09288461
| Patent No. 6159694
| GENERAL INPORMATION:
| APPLICANT: Karras, James G.
| TITLE OF INVENTION: Expression
| TITLE OF INVENTION: Expression
| TITLE OF INVENTION: Expression
| TITLE OF INVENTION: UNDER: US/09/288,461
| CURRENT APPLICATION NUMBER: US/09/288,461
| CURRENT FILING DATE: 1999-04-08
| NUMBER OF SEQ ID NOS: 107
| SGFTWARE: Patentin Ver. 2.0
| SEQ ID NO 68
| LENGTH: 20
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Pred. No. 1.2e+02;
0; Mismatches 3; Indels
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1.0%; Score 14.2; D.
Best Local Similarity 84.2%; Pred. No. 1.2e
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                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Synthetic Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 52, Application US/09488671A Patent No. 6187545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1256 GAGGCCAGGTTGAGGCCCT 1274
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       512 TCAGCGCCAACCTGCCGGA 530
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                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best_Local Similarity
Matches 16; Conserv
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US-09-488-671-52/c
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RESULT 74

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W3-09-920-672-52/c

| Sequence 52, Application US/09920672
| Sequence 52, Application US/09920672
| Sequence 52, Application US/09920672
| Patent No. 6455308
| General Information Application
| APPLICANT: Mark J. Graham
| APPLICANT: Susan M. Freier
| TITLE OF INVENTION: ANTISENSE MODULATION OF SERUM AMYLOID A4 EXPRESSION
| FILE REPERENCE: RTS-0251
| CURRENT APPLICATION NUMBER: US/09/920,672
| CURRENT FILING DATE: 2001-08-01
| SEQ ID NOS: 89
| LENGTH: 20
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                                  Score 14.2; DB 1; Length 20;
Pred. No. 1.2e+02;
0; Mismatches 3; Indels
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APPLICANT: MICHEL M. Neff

APPLICANT: MICHEL M. Neff

TITLE OF INVENTION: GENETICALLY MODIFIED PLANTS HAVING

TITLE OF INVENTION: MODULATED BRASSINGSTEROID SIGNALING

TITLE OF INVENTION: MODULATED BRASSINGSTEROID SIGNALING

TITLE OF INVENTION: MODIFIED BRASSINGSTEROID SIGNALING

TITLE OF INVENTION: MODIFIED BRASSINGSTEROID SIGNALING

TITLE OF INVENTION: 050-03-16

PRICH RELING DATE: 1999-03-16

PRICH APPLICATION NUMBER: US 60/170,931

PRICH FILING DATE: 1999-12-14

PRICH FILING DATE: 1999-12-20

NUMBER OF SEQ ID NOS: 16

SOFTHARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 4

LUNCTH: 20
                                                                                                  3; Indels
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1.0%; Score 14.2; DB 1;
Best Local Similarity 84.2%; Pred. No. 1.2e+02;
Matches 16; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Oligonuclectide primer US-09-527-073-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1037 CTGACTCTTCCCACGACAG 1055
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Patent No. 6534313
GENERAL INFORMATION:
                                                                                                                                                                      227 CTCAGCCTCAGGCATCTGC 245
                                                                                                                                                                                                        20 CTCAGCCACGGTCATCTGC 2
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                                     1.0%;
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ORGANISM: Artificial Sequence
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                                  Query Match
Best Local Similarity 84.2
Matches 16; Conservative
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Matches 16; Conserv
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US-09-851-062-47/c
Sequence 47, Application US/09851062
Facent No. 6448081
GENERAL INFORMATION:
APPLICANT: Brenda F. Baker
TITLE OF INVENTION: ANTISENSE MODULATION OF INTERLEUKIN 12 P40 SUBUNIT EXPRESSION
CURRENT APPLICATION NUMBER: US/09/851,062
CURRENT FILING DATE: 2001-05-07
NUMBER OF SEQ ID NOS: 87
LENGTH: 20
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| GENERAL INFORMATION:
| APPLICANT: Aitman, Timothy
| APPLICANT: Scott, James
| APPLICANT: Stanton, Lawrence
| TITLE OF INVENTION: Compositions and Methods of Disease Diagnosis and
| TITLE OF INVENTION: Therapy
| TITLE OF INVENTION: Therapy
| TITLE OF INVENTION THERES: 1999-03-17
| CURRENT APPLICATION NUMBER: 09/221,222
| BARLIER PILING DATE: 1999-12-23
| NUMBER OF SEQ ID NOS: 207
| SOFTWARE: PATENTIN Ver. 2.0
| SEQ ID NO 155
| LEAST OF THE 
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Pred. No. 1.2e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
1.0%; Score 14.2; DB 1;
Best Local Similarity 84.2%; Pred. No. 1.2e+02;
Matches 16; Conservative 0; Mismatches 3;
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CURRENT APPLICATION NUMBER: US/09/130,616C CURRENT FILING DATE: 1998-08-07
EARLIER APPLICATION NUMBER: 08/910,629
EARLIER FILING DATE: 1997-08-03
NUMBER OF SEQ ID NOS: 178
SEQ ID NO 123
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Synthetic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 155, Application US/09270542; Patent No. 6322976
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                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match
Best Local Similarity 84.28;
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GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: C. Frank Bennett
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INTERFERON GAMMA RECEPTOR 1 EXPRESSION
FILE REFERENCE: RTS-0234
CURRENT APPLICATION NUMBER: US/09/843,376
CURRENT FILING DATE: 2001-04-26
NUMBER OF SEQ ID NOS: 88
SEQ ID NO 62
LENGTH: 20
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APPLICANT: Tournier-Lasserve, Elisabeth
APPLICANT: Tournier-Lasserve, Elisabeth
APPLICANT: Joutel, Anne
APPLICANT: Bach, Marie-Germaine
APPLICANT: Bach, Jean-Francois
TITLE OF INVENTION: GENE INVOLVED IN CADASIL, METHOD OF DIAGNOSIS AND
TITLE OF INVENTION: TERRAPEUTICATION
TITLE OF INVENTION: TERRAPEUTICA APPLICATION
FILE REFERENCE: 03715.0048-00000
CURRANT APPLICATION NUMBER: ER 96 09733
EARLIER FILING DATE: 1999-05-11
EARLIER FILING DATE: 1997-04-16
EARLIER FILING DATE: 1997-04-16
EARLIER FILING DATE: 1997-07-31
NUMBER OF SEQ ID NOS: 163
SEQ ID NO 103
LENGTH: 20
LENGTH: 20
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7 OTHER INFORMATION: Description of Artificial Sequence: primer US-09-230-652-103
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1.0%; Score 14.2; DB 1;
Best Local Similarity 84.2%; Pred. No. 1.2e+02;
Matches 16; Conservative 0; Mismatches 3;
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                                                                                                                                                                                US-09-230-652-103/c
; Sequence 103, Application US/09230652A
; Patent No. 6537775
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                           1232 CTTTGGTGCTGGACGTGGC 1250
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                                                                       20 cirriccierredaacedec
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Best Local Similarity
Matches 16; Conserv
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US-09-422-978-11618/c
i Select 1618, Application US/09422978
Factor 10. 6537751
GENERAL INFORMATION:
APPLICANT: COhem, Daniel
APPLICANT: Chem, Daniel
APPLICANT: Chemakov, 11ya
TITLE OF INVENTION: Biallelic markers for use in constructing a high density...
TITLE OF INVENTION Biallelic markers for use in constructing a high density...
FILE REFERENCE: GENSTY.020CP1
CURRENT FILING DATE: 1999-10-20
EARLIER FILING DATE: 1999-04-21
EARLIER FILING DATE: 1998-11-23
EARLIER FILING DATE: 1998-11-23
EARLIER FILING DATE: 1998-11-23
EARLIER FILING DATE: 1998-04-21
                                                                                                                                                                                             APPLICANT: Solution Daniel
APPLICANT: Blumenfeld, Marta
APPLICANT: Chumakov, Ilya
TTULE OF INVENTION: Biallelic markers for use in constructing a high density...
FILE REFERENCE: GENSUET.02CP1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER PLING DATE: 1999-11-23
EARLIER FILING DATE: 1998-04-21
EARLIER FILING DATE: 1998-04-21
NUMBER OF SEQ ID NOS: 11796
SEQ ID NO 6216
LENGTH: 20
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; OTHER INFORMATION: upstream amplification primer 99-10151 for SEQ 2282,
US-09-422-978-6216
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                                                                                                                         Sequence 6216, Application US/09422978
Patent No. 6537751
GENERAL INFORMATION:
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19 CTCACACTTTCCACGACAG 1
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Best Local Similarity 84.2
Matches 16; Conservative
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ORGANISM: Homo Sapiens
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ORGANISM: Homo Sapiens
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LOCATION: 1..20
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                                                                       RESULT 81
US-09-422-978-6216/c
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LOCATION:
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GENERAL INFORMATION:
APPLICANT: Matteucci, Mark D.
APPLICANT: Krawczyk, Steven
TITLE OF INVENTION: SEQUENCE-SPECIFIC NONPHOTOACTIVATED
TITLE OF INVENTION: DUPLEX DNA
TITLE OF INVENTION: DUPLEX DNA
NUMBER OF SEQUENCES: 158
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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                                                                   Sequence 6, Application US/08679529
Patent No. 6171779
GENERAL INFORMATION:
APPLICANT: Chada, Kirin K.
APPLICANT: Tkachenko, Alex
APPLICANT: Tkachenko, Alex
APPLICANT: Tkachenko, Alex
APPLICANT: The Change of the Control of the Con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USAN
ZIP: 07901
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OFEWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,529
FILING DATE: 12-UL-1996
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: MACCINO, RICHART 32,538
REFERENCE/DOCKET NUMBER: UMD1-037
TELEPHONE: (908) 273-4988
TELEPAX: (908) 273-4679
INFORMATION OR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSE: Morrison & Foerster
STREET: A54 Midlefield Road, Suite 200
CITY: Menlo Park
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Richard R. Muccino
STREET: 758 Springfield Avenue
CITY: Summit
STATE: Now JETSEY
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA (genomic)
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          282 GGAAGCAGCAGCAA 295
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 19 base pairs
TYPE: nucleic acid
GTRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   unknown
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MOLECULE TYPE: DN
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCT-US91-03680-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-679-529-6
                                          US-08-679-529-6
RESULT 85
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ADDRESSEE: No. 58378470 No. 5837847disk of No. 5837847th America, Inc. STREET: 405 Lexington Avenue, 64th Floor
CITY: New York
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APPLICANT: Royer, John C
APPLICANT: Moyer, Donna L
APPLICANT: Soler, Wendy T
APPLICANT: Shuster, Jeffrey R
APPLICANT: Poll INVENTION: NON-TOXIC, NON-TOXIGENIC
TITLE OF INVENTION: FUSARIUM EXPRESSION SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%; Score 14; DB 1; Length 19;
100.0%; Pred. No. 1.1e+02;
ive 0; Mismatches 0; Indels
SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: 15 -- Amod base= OTHER OTHER INFORMATION: /mod base= OTHER OTHER INFORMATION: /note= "5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: /mod_base= OTHER
OTHER INFORMATION: /note= "5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: 18
OTHER INFORMATION: /mod_base= OTHER
OTHER INFORMATION: /note= "5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: modified_base
LOCATION: 19
OTHER INFORMATION: /mod_base= OTHER
OTHER INFORMATION: /note= "1,3-propanediol"
                                                                                                                                                                                                    4610-0011.40
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                                                 PCT/US91/03680
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                                                                                                                                                                                    REFERENCE/DOCKET UNDMEER: 4610
TELECOMMUNICATION INFORMATION:
TELERAK: 415-327-7250
TELERAK: 415-327-7250
TELERAK: 706141
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single
TOPOLOGY: linear
TOPOLOGY: linear
                                                                                                                                                                       29,959
                                              APPLICATION NUMBER: PCT/US
FILING DATE: 19910524
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MUTASHIGE, KALE H.
REGISTRATION NUMBER: 29,95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1143 CTTTTTTTTTTT 1156
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.0
Matches 14; Conservative
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CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 58378
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4592.210-US
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NAME: Starnes, Robert L.
REGISTRATION NUMBER: 41,324
REFRENCE/DOCKET NUMBER: 4592
TELECOMMUNICATION INFORMATION:
TELEPAX: 212-867-0123
TELEFAX: 212-878-9555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 2, Application US/09405564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 405 Lexington Avenue CITY: New York
TELECOMMUNICATION INFORMATION:
                     TELEPHONE: 212-867-0123
TELEPAX: 212-869-9655
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      387 AGAGGTGGCAGCAA 400
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 212-878-9655
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                            LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                 linear
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                                                                                                                                                                            ; TOPOLOGY:
US-08-816-915-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-816-239-2
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US-09-405-564-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 89
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Royer, John C
APPLICANT: Royer, John C
APPLICANT: Nover, Donna L
APPLICANT: Shuster, Jeffrey R
TITLE OF INVENTION: NON-TOXIC, NON-PATHOGENIC
TITLE OF INVENTION: FUGARIUM EXPRESSION SYSTEM
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 60603050 No. 6060305disk of No. 6060305th Am
STREET: 405 Lexington Avenue, 64th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
1.0%; Score 14; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.3e+02;
Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/921,426
FILING DATE: 29-AUG-1997
CLASSIFICATION NUMBER: US/08/921,426
FILING DATE: 29-AUG-1997
CLASSIFICATION NUMBER: US/08/456,433
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/456,433
FILING DATE: 01-JUN-1995
APPLICATION NUMBER: US 08/404,678
FILING DATE: 15-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: AGTIS DY., CHATYI H.
REGISTRATION NUMBER: 34,086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/816,915
FILING DATE: 13-MAR-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .TTORNEY/AGENT INFORMATION:
NAME: AGTIS Dr., Cheryl H.
REGISTRATION NUMBER: 34,086
REFERENCE/DOCKET NUMBER: 4216.240-US
                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 34,086
REFREENCE/DOCKET NUMBER: 4216.010-US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-867-0123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 10.74-6401
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 88
US-08-816-915-14
; Sequence 14, Application US/08816915
; Patent No. 6060305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             387 AGAGGTGGCAGCAA 400
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 212-878-9655
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid

STRANDEDNESS: single

; TOPOLOGY: linear

US-08-921-426-14
                   COUNTRY: USA
ZIP: 10174-6401
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New York
New York
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                                                   Gaps
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Patent No. 6066493
GENERAL INFORMATION:
APPLICANT: Shuster, Jeffrey R.
TITLE OF INVENTION: Morphological Mutants of Filamentous NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 1.0%; Score 14; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 1.3e+02; Matches 14; Conservative 0; Mismatches 0; Indels
  Length 20;
                                                Indels
Query Match
1.0%; Score 14; DB 1; Le:
Best Local Similarity 100.0%; Pred. No. 1.3e+02;
Matches 14; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/816,239
FILING DATE: 13-MAR-1997
CLASSIFICATION: 435
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US-09-422-978-7294/c

i Sequence 7284, Application US/09422978

i Sequence 7284, Application US/09422978

i Patent No. 6537751

i GENERAL INFORMATION:

APPLICANT: Cohen, Daniel

APPLICANT: Chumakov, Ilya

ITILE OF INVENTION: Biallelic markers for use in constructing a high density...

FILE REFERENCE: GENSET.020CP1

CURRENT FILING DATE: 1999-10-20

i BARLIER APPLICATION NUMBER: US 60/109,732

EARLIER FILING DATE: 1998-04-21

EARLIER FILING DATE: 1998-04-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COTHER INFORMATION: upstream amplification primer 99-3483 for SEQ 3360, US-09-422-978-7294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-705-390-2

Sequence 2, Application US/09705390

Sequence 2, Application US/09705390

Sequence 2, Application US/09705390

Sequence 2, Application US/09705390

SERVERAL NO. 654474

SERVERAL NO. 654474

TITLE OF INVENTION: Morphological Mutants of Filamentous TITLE OF INVENTION: Pungl
FILE REFERENCE: 4592.230-US

CURRENT APPLICATION NUMBER: US/09/705,390

CURRENT APPLICATION NUMBER: 00/010238

PRIOR APPLICATION NUMBER: 08/026114

PRIOR FILING DATE: 1996-01-19

PRIOR FILING DATE: 1996-10-04

PRIOR FILING DATE: 1996-10-04

PRIOR FILING DATE: 1997-03-13

NUMBER OF SEQ ID NOS: 2

SOFTWARE: PSELEGE for Windows Version 4.0

LENGTH: 20
  1.0%; Score 14; DB 1; Length 20; 100.0%; Pred. No. 1.3e+02; ive 0; Mismatches 0; Indels
        Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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: Pred. No. 1.3e+02;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 1.0%; Soc
Best Local Similarity 100.0%; P:
Matches 14; Conservative 0;
                                                                                                                269 GGCTGATCAAAGAG 282
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US-09-705-390-2
                                                                                                                                                                      20 GGCTGATCAAGAG 7
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                                                           14; Conservative
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Query Match
Best Local Similarity
Matches 14; Conserva
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                                                                                                                                                                                           No. 61840260 No. 6184026disk of No. 6184026th America, Inc.
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                 GENERAL INFORMATION:
APPLICANT: Shuster, Jeffrey R.
APPLICANT: Royer, John C.
TITLE OF INVENTION: Morphological Mutants of Filamentous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.0%; Score 14; DB 1; Length 20; 00.0%; Pred. No. 1.3e+02; ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                     COUNTY: USA
ZIP: 10174
COMPUTER READABLE FORM:
MEDLUM TYPE: Diskette
COMPUTER: Diskette
COMPUTER: Diskette
COMPUTER: DISKETS
SOFTWARE: FASTSEN DOS
SOFTWARE: FASTSEN OF WINDOWS Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/405,564
FILING DATE:
RISHER APPLICATION DATA:
APPLICATION NUMBER: US/08/816,239
FILING DATE: 13-MRA-1997
ATTORNY ABTORNATION:
NAME: Starnes, Robert L.
REGISTRATION NUMBER: 41,324
REFERENCE/DOCKET NUMBER: 4592.210-US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Prusiner, Stanley
APPLICANT: Tremblay, Patrick
APPLICANT: Moore, Richard
APPLICANT: Westaway, David
APPLICANT: Wood, Leroy B.
APPLICANT: Hood, Leroy B.
APPLICANT: Hoe, Inyoul
TITLE OF INVENTION: PrP-like Gene
FILE REFERENCE: 6110-130US1
CURRENT APPLICATION NUMBER: US/09/309,317
CURRENT APPLICATION NUMBER: US/09/309,317
CURRENT FILING DATE: 1999-05-11
NUMBER OF SEQ ID NOS: 21
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 7, Application US/09309317
; Patent No. 6277970
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                                                                                                                                                                                                                  STREET: 405 Lexington Avenue CITY: New York
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SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
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STRANDEDNESS: single
                                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 61840
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CORGANISM: homosapien
US-09-309-317-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                            USA
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IITLE OF INVENTION: THERMOPHILIC STRAND DISPLACEMENT AMPLIFICATION
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                   NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSE: Richard J. Rodrick, Becton Dickinson and
ADDRESSE: Company
STREET: 1 Becton Drive
                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IDN PC compatible

COMPUTER: Eloppy disk

COMPUTER: Date PC compatible

COMPANDE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/531,747

FILING DATE:

CLASSIFICATION: 435

ATTORNEY/ACENT INPORMATION:

NAME: Fugit, Donna R.

REFERATION NUMBER: 32,135

REFERATION NUMBER: 32,135

REFERATION FOR SEQ ID NO: 4:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Draper, Kenneth
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: MrSwiggen, James
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/373,124A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2029, Application US/08373124A Patent No. 5646042
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/245,466
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Fifth Street
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
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                                                                                                                        Franklin Lakes
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STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                 TRY: US
07417
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                                                                                                                        CITY: Frank
STATE: NJ
                                                                                                                                                                    COUNTRY:
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GENERAL INFORMATION:
APPLICANT:
APPLICANT:
TITLE OF INVENTION: FUSARIUM EXPRESSION SYSTEM AND PROMOTERS AND TERMINATORS FOR CORRESPONDENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nowo Nordisk of North America, Inc.
STREET: 405 Lexington Avenue, 64th Floor
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Patent No. 5631147
GENERAL INFORMATION:
APPLICANT: Lohman, Kenton L.
APPLICANT: Van Clerova, Matalie V.
APPLICANT: Reid, Robert A.
TITLE OF INVENTION: DETECTION OF NUCLEIC ACIDS IN CELLS BY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.0%; Score 14; DB 1; Length 20;
L00.0%; Pred. No. 1.38+02;
      Score 14; DB 1; Length 20;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/07743
FILING DATE: 15-June-1995
                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Novo Nordisk of North America, Inc. STREET: 405 Lexington Avenue, 64th Floor CITY: New York STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 100.0%; Pred. No. 1.3
Matches 14; Conservative 0; Mismatches
    Query Match 1.0%; Score 14; DB Best Local Similarity 100.0%; Pred. No. 1.3 Matches 14; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 10174-6401
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                387 AGAGGIGGCAGCAA 400
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EDNESS: single
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US-08-531-747-4/c
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PCT-US95-07743-14
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/531,747
                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/08781432
Patent No. 5756702
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA (genomic) US-08-781-432-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17 Archargaggaagcrec 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Fugit, Donna R. REGISTRATION NUMBER: 32, REFERENCE/DOCKET NUMBER: INFORMATION FOR SEQ ID NO: 4 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               274 ATCAAAGAGGAAGCAGC
                                                  Query Match
Best Local Similarity 88.23
Matches 15; Conservative
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Best Local Similarity 88.2
Matches 15; Conservative
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07417
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STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 99
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APPLICANT: Lohman, Kenton L.
APPLICANT: Ostrerova, Natalie V.
APPLICANT: Van Cleve, Mark
APPLICANT: Reid, Rober, Mark
TITLE OF INVENTION: DETECTION OF NUCLEIC ACIDS IN CELLS BY
TITLE OF INVENTION: THERMOPHILIC STRAND DISPLACEMENT AMPLIFICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 17;
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Richard J. Rodrick, Becton Dickinson and
ADDRESSEE: Company
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 07417
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: BHE PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/531,749
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.0%; Score 13.8; E
47.1%; Pred. No. 97;
tive 7; Mismatches
                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 229/035
REFERENCE DOCKET NUMBER: 209/035
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: 67-3510
INFORMATION FOR SEQ ID NO: 2029:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: MUCHOL CARACTERISTICS:
TYPE: M
                                                                  FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
FILING DATE: August 26, 1992
ATONNEY/AGENT INFORMATION:
NAME: WArburg, Richard
REGISTRATION NUMBER: 32,327
May 18, 1922, 943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Fugit, Donna R.
REGISTRATION NUMBER: 32,135
REFERENCE/DOCKET NUMBER: P-34
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4, Application US/08531749; Patent No. 5733752; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1094 TIGAACGIAATTAIGIA 1110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 UUGAAAGUUAUUAUGUA 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 Becton Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Franklin Lakes
                    FILING DATE: May 18
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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Best Local Similarity
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US-08-373-124A-2029
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US-08-531-749-4/C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-531-749-4
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APPLICANT: Lohman, Kenton L.
APPLICANT: Ostracova, Matalie V.
APPLICANT: Van Cleve, Mark
APPLICANT: Van Cleve, Mark
APPLICANT: Reid, Robert A.
TITLE OF INVENTION: DETECTION OF NUCLEIC ACIDS IN CELLS BY
TITLE OF INVENTION: THERMOPHILIC STRAND DISPLACEMENT AMPLIFICATION
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
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  DB 1; Length 17;
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                                             2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/781,432
1.0%; Score 13.8; D
88.2%; Pred. No. 97;
iive 0; Mismatches
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88.2%; Pred. No. 97;
ive 0; Mismatches
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Patent No. 5817796
GENERAL INFORMATION:
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Fatent No. 6057156
GENERAL INFORMATION:
APPLICANT: Aktar, Saghir
APPLICANT: Aktar, Saghir
APPLICANT: McSwiggen, James
TITLE OF INVENTION: OF DISEASES OR CONDITIONS RELATED
TITLE OF INVENTION: TO LEYELS OF EPIDERMAL GROWTH
TITLE OF INVENTION: PACTOR RECEPTORS
NUMBER OF INVENTION: FACTOR RECEPTORS
NUMBER OF SEQUENCES: 1877
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: 631 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                               COUNTY: CALLLY...

COUNTY: U.S.A.

ZIP: 90071-2066

CMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: Enstsed for Windows 2.0
SOFTWARE: FRASER for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 08/08/985,162
FILING DATE: 04 December 1997
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/036,476
FILING DATE: 31 January 1997
APPLICATION NUMBER: 230/107
FILENCHERNCE,DOCKET NUMBER: 230/107
TELECOMONICATION INFORMATION:
TELECOMONICATION TREADMATION:
TELECOMONICATION TREADMATION TREADM
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FASTENG for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/985,162
      633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               629 AGCTCCAGGAGCTCTGC 645
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Best Local Similarity 88.2:
Matches 15; Conservative
                                          Suite 4700
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STATE: California
COUNTRY: U.S.A.
                                                                     Los Angeles
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JS-08-985-162-645/c
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      STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17, Application US/08985162
Patent No. 6057156
GENERAL INFORMATION:
APPLICANT: Akhtar, Saghir
APPLICANT: Fell, Patricia
APPLICANT: McSwiggen, James
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT
TITLE OF INVENTION: OF DISEASES OR CONDITIONS RELATED
TITLE OF INVENTION: TO LEVELS OF EPIDERMAL GROWTH
TITLE OF INVENTION: PACTOR RECEPTORS
METHODS AND COMPOSITIONS FOR
TREATMENT OF RESTENOSIS AND
CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: MAND PETECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,628
FILING DATE: 05-MAY-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/373,124
FILING DATE: MAND 18, 1995
APPLICATION NUMBER: 08/245,466
FILING DATE: MAY 18, 1994
APPLICATION NUMBER: 08/373,132
FILING DATE: PEDUARY 7, 1994
APPLICATION NUMBER: 07/98/132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/98/132
FILING DATE: ADMINISTER: 07/98/132
FILING DATE: ADMINISTER: 07/98/132
FILING DATE: ADMINISTER: 07/98/132
FILING DATE: ADMINISTER: 07/98/132
APPLICATION NUMBER: 07/98/132
FILING DATE: AUGUST 26, 1992
ATTONEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard
REISTRATION NUMBER: 32,327
REPRENCE/POCKET NUMBER: 209/035
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                        E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1094 TTGAACGTAATTATGTA 1110
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1 UUGAAAGUUAUUAUGUA 17
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 67-3510
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lyon & Lyon
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Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                     NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Ly
                                                                                                                                                                                                                                                 CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071
                                                                                                                                                                                                                          Suite 4700
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 100
US-08-985-162-17/c
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US-08-435-628-2029
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                                                                                                                                                                                     STREET:
STREET:
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Patent No. 5877021

GENERAL INFORMATION:
APPLICANT: Stinchcomb, Daniel T.
APPLICANT: MICHORAL JAMES
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
NUMBER OF SEQUENCES: 2751
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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88.2%; Pred. No. 97;
tive 0; Mismatches
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PRIOR FILING DATE: 1997-11-05
PRIOR FILING DATE: 1997-11-05
PRIOR FILING DATE: 1997-11-05
PRIOR APPLICATION NUMBER: US 60/084,727
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR APPLICATION NUMBER: US 09/301,511
PRIOR APPLICATION NUMBER: US 09/301,511
PRIOR FILING DATE: 1999-04-28
NUMBER OF SEQ ID NOS: 1526
SOFTWARE: PATCHIN VERSION 3.0
SEQ ID NO 684
                                                                                                                                                                                                                   Patent No. 6528640
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 90071
COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                Sequence 684, Application US/09474432B Patent No. 6528640
                                                   274 ATCAAAGAGGAAGCAGC 290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             517 GCCAACCTGCCGGAGGA 533
                                                                                            17 ATCAATGAGGAAGCTGC
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                               APPLICANT: Beigelman, Leo
APPLICANT: Burgin, Alex
APPLICANT: Beaudry, Amber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-585-684B-2548/c
                                                                                                                                                       RESULT 103
US-09-474-432B-684
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88.2%; Pred. No. 97;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.0%; Score 13.8; DB 1; Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Richard J. Rodrick - Becton, Dickinson and ADDRESSEE: Company STREET: 1 Becton Drive CITY: Franklin Lakes STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 07417
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/964,020
FILING DATE:
CLASSIFICATION 514

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/036,476
FILING DATE: 31 January 1997
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 230/107
TELECHMUNICATION INFORMATION:
TELECHMUNICATION INFORMATION:
TELECHMUNICATION 1999-1600
TELECHARM: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Higher, David W.
REGISTRATION NUMBER: 30,265
REFERENCE/DOCKET NUMBER: p-402
TELECOMMUNICATION INFORMATION:
TELEPHONE: (201) 847-5317
TELEPAX: (201) 846-9228
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                       TELERX: (213)
TELERX: 67-3510
INFORMATION FOR SEQ ID NO: 645:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         916 CTAAAGGAGATGGCAGA 932
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 15; Conserva
                                                                                                                                                                                                                                                                                                                                                         ; TOPOLOGY: linear
US-08-985-162-645
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-964-020-2
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APPLICANT: Beaudry, Amber
APPLICANT: Rarpeisky, Alex
APPLICANT: Rarpeisky, Alex
APPLICANT: Adamic, Jasenka
APPLICANT: Sweedler, David
APPLICANT: Sweedler, David
APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleotide triphosphate and their incorporation into oligonuclee
FILE REPERENCE: MEHROD-831-B (247/276)
CURRENT APPLICATION NUMBER: US/09/474,432B
CURRENT FILING DATE: 1999-12-19
                                   0
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     88.2%; Pred. No. 97; tive 0; Mismatches
     Best Local Similarity 88.2
Matches 15; Conservative
04 December 1997
N: 514
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INFORMATION FOR SEQ ID NO: 33.
SEQUENCE CHARACTERISTICS.
LENGTH. 18 base pairs
TYPE: mucleic acid
STRANDENESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        464 GCAGCCTGCAGGGGAG 480
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                                (202) 408-4400
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                      Local Similarity 88.2 es 15; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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: USA
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                                TELEFAX:
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Matches
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Sequence 33. Application US/08702105A

Patent No. 5908839

GENERAL INFORMATION

APPLICANT: Levitt, Roy C.

APPLICANT: Maloy, W. Lee

APPLICANT: Maloy, W. Lee

APPLICANT: Nicolaides, Nicholaides, Nicholaides, Nicholaides, Nicholaides, TITLE OF INVENTION: Treating Acopic Allergies Including Asthma And Related

TITLE OF INVENTION: Treating Acopic Allergies Including Asthma And Related

TITLE OF INVENTION: Disorders

NUMBER OF SEQUENCES: 41

CORRESPONDENCE ADDRESS:

ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & ADDRESSEE: Dunner L.L.P.

STREET: 1300 I Street N.W., Suite 700

CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
FILING DATE: US/08/702,105A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 05387.0056-01000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,684B
FILLNG DATE: January 16, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/000,951
FILLNG DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION 1000 155-0440
TELEPAX: (213) 955-0440
TELEPAX: (213) 955-0440
INFORMATION FOR SEQ ID NO: 2548:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: D.C.
COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/874,503
FILING DATE: 13-UUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: FORGIS, Jean B.
REGISTRATION NUMBER: 32984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1157 GGAAGTAAAGCAGCTAA 1173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18 GGAAGCAAAGCAGGTAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TOPOLOGY: linear
US-08-585-684B-2548
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
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US-08-702-105A-33/c
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US-08-702-110A-33/C
; Sequence 33, Application US/08702110A
; Sequence 33, Application US/08702110A
; Patent No. 6037149
; GENERAL INFORMATION:
APPLICANT: Maloy, W. Lee
APPLICANT: Maloy, W. Lee
APPLICANT: Maloy, W. Lee
APPLICANT: Nicolaides, Nicholas C.
TITLE OF INVENTION: Treating Asthma Associated Factors As Targets For
TITLE OF INVENTION: Treating Atopic Allergies Including Asthma And Related
TITLE OF INVENTION: Disorders
NUMBER OF SEQUENCES:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
ADDRESSEE: Dunner L.L.P.
STREET: 1300 I Street N.W., Suite 700
CITY: Washington ó · 0 Gaps Gaps ö . 1.0%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 1.1e+02; Length 18; Indels Indels ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/702,110A
FILING DATE: Score 13.8; DB 1; Pred. No. 1.1e+02; 0; Mismatches 2; CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION DATA:
APPLICATION UNBER:
PULING DATE: 13-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Fordis, Jean B.
REGISTRATION NUMBER: 32984
REFERENCE/DOCKET NUMBER: 05387.0056-01000
TELECOMMUNICATION INFORMATION:
TELEFAX: (202) 408-4000
TELEFAX: (202) 408-4000 0; Mismatches

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| US-09-630-706-86/C | US-09630706 |
| Sequence 86, Application US/09630706 |
| Patent No. 6277640 |
| GENERAL INFORMATION: |
| APPLICANT: C. Frank Bennett |
| APPLICANT: Lex M. Cowsert |
| TITLE OF INVENTION: ANTISENSE MODULATION OF HER-3 EXPRESSION |
| TITLE REFERENCE: RTS-0063 |
| CURRENT APPLICATION NUMBER: US/09/630,706 |
| CURRENT FILING DATE: 2000-08-01 |
| NUMBER OF SEQ ID NOS: 94 |
| LENGTH: 18 |
| LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.0%; Score 13.8; DB 1; Length 18;
88.2%; Pred. No. 1.1e+02;
tive 0; Mismatches 2; Indels
                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
ADDRESSEE: Dunner L.L.P.
                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/325,571
                                                                                                                                                                                                                                                                                                                                                                          FILING DAIL:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/874,503
FILING DATE: 13-UN-1997
ATTORNEY/AGENT INFORMATION:
NAME: FORGIS, Jean B.
REGISTRATION NUMBER: 32984
REFERENCE/DOCKET NUMBER: 05387.0056-01000
TELECOMMUNICATION INFORMATION:
TELEFRAN: (202) 408-4000
TELEFAX: (202) 408-4000
TELEFAX: (202) 408-4100
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 1.0%; Score 13.8; DB 1; Best Local Similarity 88.2%; Pred. No. 1.1e+02; Matches 15; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ), OTHER INFORMATION: Antisense Oligonucleotide US-09-630-706-86
                                                                                              %: 1300 I Street N.W., Suite 700 Washington
                                                                                                                                                              COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE PORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       464 GCAGCCTGCAGGGGAG 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        977 CTTGACCAGTCCCATTC 993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17 GTAGGCTGCAGGGGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18 CTTTCCCAGTCCCATTC
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Matches 15; Conserv
                                                                                              STREET: 130
CITY: Washin
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-325-571-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          à
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APPLICANT: Levitt, Roy C.
APPLICANT: Maloy, W. Lee
APPLICANT: Kari, U. Prasad
APPLICANT: Kari, U. Prasad
APPLICANT: Nicolaides, Nicholas C.
TITLE OF INVENTION: Asthma Associated Factors As Targets For
TITLE OF INVENTION: Treating Atopic Allergies Including Asthma And Related
TITLE OF INVENTION: Disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
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                                                                                                             US-09-038-073-2548/c
; Sequence 2548, Application US/09038073
; Patent No. 6194150
; GENERAL INFORMATION:
APPLICANT: Stinchcomb, Daniel T.
APPLICANT: McSwigen, James
; APPLICANT: McSwigen, James
; TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
; TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
; TITLE OF SEQUENCES: 2751
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 1.0%; Score 13.8; DB 1; Best Local Smilarity 88.2%; Pred. No. 1.1e+02; Matches 15; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPOTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/038,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       218/078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION UNDBR: 32,327
REFERENCE/DOCKET NUMBER: 218/07
FELECOMMUNICATION INFORMATION:
TELERAX: (213) 489-1600
TELERAX: (713) 955-0440
INFORMATION FOR SEQ ID NO: 2548:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 108
US-09-325-571-33/c
; Sequence 33, Application US/09325571
; Patent No. 6261559
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/585,684
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1157 GGAAGTAAAGCAGCTAA 1173
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17 GTAGGCTGCAGGGGAG 1
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EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: linear
US-09-038-073-2548
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A. ZIP: 90071
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Gaps

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Gaps

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Indels

Length 18;

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RESULT 112
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                                                                                                                                                     APPLICANT: Merlo, Parricia Ann Owens
APPLICANT: Guo, Lining
APPLICANT: Guo, Lining
APPLICANT: Scokut, Thomas A.
APPLICANT: Young, Scott A.
APPLICANT: Young, Scott A.
APPLICANT: Folkerts, Otto
APPLICANT: Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR
TITLE OF INVENTION: IN PLANTS
NUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: UJ1y 12, 1996
CLASSIFICATION: 800
PRIOR APPLICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/001,135
RILING DATE: UJ1y 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: SEPTEMBER: 0,1994
ATTORNEY AGENT INFORMATION:
US-001-679-645-583
; Sequence 583, Application US/08679645
; Patent No. 6350934
; GENERAL INFORMATION:
; APPLICANT: Zwick, Michael G.
; APPLICANT: Edington, Brent E.
; APPLICANT: McSwiggen, James A.
; APPLICANT: McSwiggen, James A.
; APPLICANT: Guo, Lining
; APPLICANT: Skokut, Thomas A.
; APPLICANT: Young, Scott A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
ZIE: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TREECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 111
US-08-535.249-98
Sequence 98, Application US/08535249
Patent No. 6455689
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                        E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 1.0%;
Best Local Similarity 70.6%;
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERICS:
LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear
US-08-679-645-583
                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
STREET:
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APPLICANT: Schlingensiepen, Georg-Ferdinand
APPLICANT: Brysch, Wolfgang
APPLICANT: Schlingensiepen, Karl-Hermann
APPLICANT: Schlingensiepen, Ramar
APPLICANT: Schlingensiepen, Reimar
APPLICANT: Scholand, Ulrich
TITLE OF INVENTION: immuno-suppressive effect of transforming-growth-factor beta
NUMBER OF SEQUENCES: 137
CORRESPONDENCE ADDRESS:
ADDRESSEE: Jacobson, Price, Holman & Christian STREET: 400 Seventh Christian STREET: AND STREET: AND STREET: AND STREET: AND STREET: AND STREET: AND ST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ·.
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TITLE OF INVENTION: Chromosomal Markers and Diagnostic Tests for Manic-Depressive Illness CORRESPONDENCES: 197
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: EC-DOS/MS-LOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/535,249
FILING DATE:
CLASSIFICATION S14
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 93 107 089.0
FILING DATE: 30-APR-1993
PRIOR APPLICATION NUMBER: BP 93 107 849.7
FILING DATE: 30-APR-1093
ATTORNEY AGENT INFORMATION:
NAME: Player, William E.
REFERENCE/POCKET NUMBER: 10577/P58418
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ.ID NO: 98:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 193, Application US/09091952A
; Patent No. 6458532
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1018 AGATGGTGCCAAAGTGC 1034
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGATGGTACAAAAGTGC
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LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Washington D.C
XY: U.S.A.
20004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-535-249-98
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us09904568-3.rni

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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 49, Application US/07741940 Patent No. 5352775
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100 ACAACCCCGGAGGCGCA 116
                                                                                                                                                                                                                                                935 TGGAGAAGAGGTGTGAG 951
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ANAND, RAKESH
CARLSON, MARY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GRODEN, JOANNA
HEDGE, PHILIP J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KINZLER, KENNETH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                        Query Match 1.0
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JOSLYN, GEOFF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: NUCLEIC ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 2001-4598
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
E: cDNA
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Wash:
STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                     US-07-741-940-49/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-07-741-940-49
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APPLICANT: Cohen, Daniel
APPLICANT: Cohen, Daniel
APPLICANT: Cohen, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Chumakov, Ilya
TITLE OF INVENTION: Biallelic markers for use in constructing a high density...
FILE REPERROR: CENERY TO20cP1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US 60/298,850
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER FILING DATE: 1998-11-23
EARLIER FILING DATE: 1998-11-23
EARLIER FILING DATE: 1998-11-30
NUMBER FILING DATE: 1998-04-21
NUMBER FILING DATE: 1998-04-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Pred. No. 1.1e+02;
0; Mismatches 2; Indels
                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette

COMPUTER IBM Compatible

COMPUTER: IBM Compatible

COMPUTER: IBM Compatible

OPERATING SYSTEM: DOS

SOFTWARE: FastSEQ for Windows Version 2.0

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/091,952A

FILING DATE: 19-Apr-1999

CLASSIFICATION NUMBER: US 60/029,278

APPLICATION NUMBER: PCT/US97/19381

FILING DATE: 28-OCT-1997

ATTORNEY/AGENT INFORMATION:

NAME: SMITH, TIMOCHAY I.

REGISTRATION NUMBER: 35,367

ATTORNEY/AGENT INFORMATION:

NAME: SMITH, TIMOCHAY I.

REGISTRATION NUMBER: 35,367
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: 1...18
OTHER INFORMATION: Clone 47 reverse primer
SEQUENCE DESCRIPTION: SEQ ID NO: 193:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 015280-297100US
TELECOMMUNICATION INFORMATION:
TELEPEONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4727, Application US/09422978
Patent No. 6537751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1319 GTGCTTTTGTAGATCTT 1335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: <UDknown>
INFORMATION FOR SEQ ID NO: 193:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Grechtererakecrehr 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 88.2%;
Matches 15; Conservative
                                          San Francisco
                                                                                          COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-091-952A-193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-422-978-4727
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LENGTH: 18
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FEATURE:
NAME/KEY: primer\_bind
LOCATION: 1..18
COTHER INPORMATION: upstream amplification primer 99-17363 for SEQ 793, US-09-422-978-4727 Gaps Gaps .. 0 ö APPLICANT: MARKANIA, ALEXANDER F.
APPLICANT: NAKANURA, YUSUKB
APPLICANT: THILVINSI NANDRA SAND SOMATIC MUTATIONS OF APC
TITLE OF INVENTION: GENE IN COLORECTAL CANCER IN HUMANS 1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 1.38+02; tive 0; Mismatches 2; Indels Length 18; Indels CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/741,940
FILING DATE: 19920109
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Kagan, Sarah A. 32,141
REGISTRATION NUMBER: 32,141
REGISTRENCE/DOCKET NUMBER: 1107.035574
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION: 1.0%; Score 13.8; DB 1; 88.2%; Pred. No. 1.1e+02; tive 0; Mismatches 2; NUMBER OF SEQUENCES: 94
CORRESPONDENCE ADDRESS:
ADDRESSEE: Barner, Birch, McKie & Beckett
STREET: 1001 G Street, NW

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APPLICANT: KINZLER, KENNETH
APPLICANT: MARKHAM, ALEXANDER F.
APPLICANT: MARKHAM, ALEXANDER F.
APPLICANT: THALFERIS, ANDREW
TITLE OF INVENTION: INHERITED AND SOMATIC MUTATIONS OF APC
TITLE OF INVENTION: GENE IN COLORECTAL CANCER IN HUMANS
NUMBER OF SEQUENCES: 94
CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner, Birch, McKie & Beckett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JOSLYN, GEOFF
KINZLEN, KENNETH
MARKHAM, ALEXANDER F.
NAKAMURA, YUSUKE
THLIVERIS, ANDREW
VERMICH INHERITED AND SOMATIC MUTATIONS OF APC
VENTION: GENE IN COLORECTAL CANCER IN HUMANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 13.8; DB 1; Length 19;
Pred. No. 1.3e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                    CITY: Mac...
STATE: D.C.
COUNTRY: USA
ZIP: 20001-4598
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
CURRART APPLICATION DATA::
CURRART APPLICATION NUMBER: US/08/452,654
PTLING DATE: 25-MAY-1995
TLING DATE: 25-MAY-1995
                                                                                                                                                                                                                                                                                                     ADDRESSEE: Banner, Birch, McKie & Beckett
STREET: 1001 G Street, NW
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1107.035574
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 25-MAY-1995
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/741,940
FILING DATE: 08-AUG-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 49, Application US/08452655B
; Patent No. 578366
; GENERAL INFORMATION:
APPLICANT: ALBERTSEN, HANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
REFRENCE/DOCKET NUMBER: 1107
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
TELEPRAX: 202-508-9299
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100 ACAACCCCGGAGGCGCA 116
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ANAND, RAKESH
CARLSON, MARY
GRODEN, JOANNA
HEDGE, PHILIP J.
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88.2%;
                             PHILIP J.
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Homo sapiens
US-08-452-654-49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-452-655B-49/C
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APPLICANT:
APPLICANT:
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INHERITED AND SOMATIC MUTATIONS OF APC
GENE IN COLORECTAL CANCER IN HUMANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE:

COUNTRY: USA

ZIP: 20001-4598

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/289,548A
FILING DATE: 12-AUG-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Kagan, Sarah
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 1107.46943
TELECOMPUNICATION INFORMATION:
TELECOMPUNICATION INFO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Banner & Allegretti, LTD
STREET: 1001 G Street, NW
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 116
US-08-452-654-49/c
US-08-452-654-49/c
; Patent No. 5691454
; GENERAL INFORMATION:
APPLICANT: ALBERTSEN, HANS
; APPLICANT: ANAND, RAKESH
; APPLICANT: CARLSON, WARX
                                                                                                                                                               Sequence 49, Application US/08289548A
Patent No. 5648212
GENERAL INFORMATION:
APPLICANT: ALBERTSEN, HANS
APPLICANT: ANAND, RAKESH
APPLICANT: CARLSON, MARY
                                                                                                                                                                                                                                                                             ANAND, RAKESH
CARLSON, MARY
GRODEN, JOANNA
HEDGE, PHILIP J.
JOSLYN, GEOFF
KINZLER, KENNETH
MARKHAM, ALEXANDER F.
NAKAMURA, YUSUKE
THLIVERIS, ANDREW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100 ACAACCCCGGAGGCGCA 116
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17 ACAACCCAGGAGCCGCA 1
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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: CDNA
PORGINAL SOURCE:
CRGANISM: HOMO sapiens
US-08-289-548A-49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: INH
TITLE OF INVENTION: GEN
UNMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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STATE: D.C.
                                                                                                             RESULT 115
US-08-289-548A-49/c
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                     APPLICANT
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Gaps
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US-08-471-973A-33/C
Sequence 33, Application US/08471973A
Sequence 33. Application US/08471973A
Patent No. 5872232
GARERAL INFORMATION:
APPLICANT: Phillip Dan Cook
APPLICANT: Andrew Kawasaki
TITLE OF INVENTION: Sugar Modified Oligonucleotides
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
ADDRESSE: Woodcock Washburn Kurtz Mackiewicz and No. 5872232ris
STREET: One Liberty Place - 46th Floor
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.0%; Score 13.8; DB 1; Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: FAA

COUNTRY: U.S.A.

ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 720 Kb
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/471,973A
FILING DATE: 06-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 335,932
FILING DATE: 05-WAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: JOSEPH LUCCI
REGISTRATION NUMBER: 33,307
REFERENCE/DOCKET NUMBER: 1SIS-2005
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
                       CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 835,932
FILING DATE: 05-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: JOSEPH LUCCI
REGISTRATION NUMBER: 33,307
REFERENCE/DOCKET NUMBER: ISIG-
TELECHONE: 215-568-3100
TELEFRAX: 215-568-3100
TELEFRAX: 215-568-3100
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1141 GCCTTTTTTTTTTT 1157
         06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
TOPOLOGY:
US-08-468-037A-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-471-973A-33
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Sequence 33, Application US/08468037A

Fatent No. 5859221

GENERAL INFORMATION:

APPLICANT: Phillip Dan Cook

APPLICANT: A. Kawasaki

TITLE OF INVENTION: 2'-Modified Oligonucleotides

NUMBER OF SEQUENCES: 37

CORRESPONDENCE ADDRESS:

ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 5859221ris

STREET: One Liberty Place - 46th Floor

CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .;
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                                                                                                                                                                                                                                                                                                   CMPUTER: IEBM PC compatible
COMPUTER: IEBM PC compatible
COMPUTER: IEBM PC compatible
COMPUTER: IEBM PC compatible
SUSTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/452,655B
FILING DATE: 25-MAY-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/289,548
FILING DATE: 12-AMG-1994
FRIUNG DATE: 12-AMG-1994
FRIUNG DATE: 08-AMG-1994
FRIUNG DATE: 08-AMG-1994
FRIUNG DATE: 08-AMG-1994
FREFERENCE/COCKET NUMBER: 1107,49964
FREFERENCE/COCKET NUMBER: 1107,49964
TELEGRAX: 202-508-9100
TELEFRAX: 202-508-9100
TELEFRAX: 202-508-9299
INFORMATION FOR EQ. ID NO: 49:
SEQUENCE CHARACTERISTICS:
LEMGTH: 19 base pairs
TWENT: THE TELEGRAY TO THE TE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 13.8; DB 1;
Pred. No. 1.3e+02;
0; Mismatches 2;
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ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 720 Kb
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDBERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/468,037A
                                                           ADDRESSEE: Banner & Witcoff, Ltd.
STREET: 1001 G Street, NW
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100 ACAACCCGGAGGCGCA 116
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                                                                                                                                                                                                          ZIP: 2001-4598
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; ORGANISM: Homo sapiens
US-08-452-655B-49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 88.28
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDNA
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                                                                                                                                                     STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: PA
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                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ð
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APPLICANT: NAKAMURA, YUSUKE
APPLICANT: THLIYURIS, ANDREW
TITLE OF INVENTION: INHERITED AND SOMATIC MUTATIONS OF APC
TITLE OF INVENTION: GENE IN COLORECTAL CANCER IN HUMANS
NUMBER OF SEQUENCES: 102
CORRESPONDENCE ADDRESS:
ADDRESSE: Banner & Witcoff, Ltd.
STREET: 1001 & Street, NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING THE COMPANDES SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/450,582 FILING DATE: CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                       ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 720 Kb
COMPUTER: ISM PC compatible
COMPUTER: ISM PC-DOS/MS-DOS
SOFTMARE: WOATGBEIFCET 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ISIS-2004
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CCUNTRY: USA
ZIP: 20001-4598
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
"""IFR: IBM PC compatible
""" PC-DOS/MS-DY
                                                                                                                                                                                                                                                                                                                                                            APPLICATION.

FILING DATE:
ATTORNAY AGENT INFORMATION:
NAME: JOSEPH LUCCI
RECISTRATION NUMBER: 33,307
REFERENCE/DOCKET NUMBER: 1SIS-
TELECOMMUNICATION INFORMATION:
TELEFAX: 215-568-3100
TELEFAX: 215-568-3439
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 bases
LENGTH: 19 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 49, Application US/08450582 Patent No. 6114124 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/468,037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KINZLER, KENNETH
MARKHAM, ALEXANDER F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1141 GCCTTTTTTTTTTG 1157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ALBERTSEN, HANS
ANAND, RAKESH
CARLSON, MARY
GRODEN, TOANNA
HEDGE, PHILIP J.
JOSLYN, GEOFF
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Best Local Similarity 88.2'
Matches 15; Conservative
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STRANDEDNESS: single
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                  U.S.A.
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TOPOLOGY:
US-09-035-357-33
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APPLICANT:
                     COUNTRY:
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APPLICANT: Phillip Dan Cook
APPLICANT: A. Kawasaki
TITLE OF INVENTION: 2'-Modified Oligonucleotides
NUMBER OF SEQUENCES: 3'
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 6005087ris
STREET: One Liberty Place - 46th Floor
CITY: Philadelphia
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1.3e+02;
-haq 2; Indels
                                             Mismatches
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PRIOR APPLICATION DATA,

PRIOR APPLICATION NUMBER: 244,993

FILING DATE: 21-JUN-1994

ATTORNEY/AGENT INFORMATION:
NAME: JOSEPH LUCCI
REGISTRATION NUMBER: 33,307

REFERENCE/DOCKET NUMBER: 151S-2002

TELECOMMUNICATION INFORMATION:
TELEFAX: 215-568-3100

INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LINGTH: 19 bases
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                     Pred. No.
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US-09-035-357-33/c
; Sequence 33, Application US/09035357
; Patent No. 6005087
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                     88.2%;
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Best Local Similarity 88.2
Matches 15; Conservative
                     Best Local Similarity 88.2
Matches 15; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: One Liberty
CITY: Philadelphia
STATE: PA
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US-08-465-880-28/c
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Score 13.8; DB 1; Length 19;
Pred. No. 1.3e+02;
0; Mismatches 2; Indels
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Gequence 4, Application US/09130973;
Facquence 4, Application US/09130973;
Facquence 4, Application US/09130973;
Facquence 4, Application US/09130973;
Facquence 4, Application US/09130973;
FAPLICANT: Manabar, Muthiah
FAPLICANT: Cook, Phillip Dan
APPLICANT: Rawasaki, Andraw M
TITLE OF INVENTION: Making Same
TITLE OF INVENTION: Making Same
FILE REPRENENCE: ISIS2955;
CURRENT APPLICATION NUMBER: US/09/130,973;
CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
SEQ ID NO 4
ILENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: No. 6172209e1
; OTHER INFORMATION: Sequence
US-09-130-973-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.0%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 1.3e+02; tive 0; Mismatches 2; Indels
                              APPLICANT: MONIA, Breat P
TITLE OF INVENTION: Gapped 2' Modified Oligonucleotides
FILE REFERENCE: 1818313
CURRENT APPLICATION NUMBER: US/09/144,611A
CURRENT FILING DATE: 1998-08-31
PRIOR PPLICATION NUMBER: 08/861,306
PRIOR FILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 12
LENGTH: 19
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OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: Sequence
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APPLICANT: Mancharan, Muthiah
APPLICANT: Kawasaki, Andrew
IITLE OF INVENTION: Aminoxy-Modified Oligonuclectides
FILE REFERENCE: ISIS2824
CURRENT APPLICATION NUMBER: US/09/477,902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 1.0%; Score 13.8; DB 1; I
Best Local Similarity 88.2%; Pred. No. 1.3e+02;
Matches 15; Conservative 0; Mismatches 2;
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Patent No. 6194598
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                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence
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Best Local Similarity 88.23
Matches 15; Conservative
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APPLICANT: Cook, P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 125
US-09-130-973-4/c
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US-09-477-902-4/c
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Sequence A Application US/09016520A

Batent No. 612753

GENERAL INFORMATION:
APPLICANT: Cook, Phillip D

APPLICANT: Manoharan, Muthiah
APPLICANT: Rawasaki, Andrew
TITLE OF INTENTION: Aminooxy-Modified Oligonucleotides
FILE REFERENCE: 151S2824

CURRENT APPLICATION NUMBER: US/09/016,520A

CURRENT APPLICATION NUMBER: 60/037,143

EARLIER FILING DATE: 1998-01-30

EARLIER FILING DATE: 1998-01-4

NUMBER OF SEQ ID NOS: 47

SOFUMBER: Patentin Ver: 2.1

LENGTH: 19
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1.0%; Score 13.8; DB 1;
Best Local Similarity 88.2%; Pred. No. 1.3e+02;
Matches 15; Conservative 0; Mismatches 2;
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Pred. No. 1.3e+02;
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APPLICATION NUMBER: US 08/452,655
FILING DATE: 25-MAY-1995
FILING DATE: 12-MOG-1994
FILING DATE: 12-MOG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/741,940
FILING DATE: 08-MG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 1107.49964
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF $20.508-9100
TELECOMMUNICATION OF $20.508-9299
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
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US-09-144-611-12/c
Sequence 12, Application US/09144611A
Patent No. 6146829
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Best Local Similarity 88.2%;
Matches 15; Conservative
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US-09-135-202-33/c
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US-09-477-902-4
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US-09-315-886C-32/C
; Sequence 32, Application US/09315886C
; Patent No. 6225063
; GENERAL INFORMATION:
; APPLICANT: Khvorova, Anastasia
; APPLICANT: Yarus, Michael
; TITLE OF INVENTION: RNA Channels in Biological Membranes
; FILE REFERENCE: UTC-03444
; CURRENT APPLICATION NUMBER: US/09/315,886C
; CURRENT FILING DATE: 1999-05-20
; PRIOR FILING DATE: 1999-05-22
; NUMBER OF SEQ ID NOS: 32
; SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                         1.0%; Score 13.8; DB 1; Length 19;
88.2%; Pred. No. 1.3e+02;
Live 0; Mismatches 2; Indels
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1.0%; Score 13.8; DB 1; Length 19;
Best Local Similarity 88.2%; Pred. No. 1.3e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
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GENERAL INFORMATION:
APPLICANT: Cook, Phillip Dan
APPLICANT: Monia, Brett P.
TITLE OF INVENTION: Gapped 2-Modified Oligonucleotides
FILE REFERENCE: ISIS-4291
CURRENT APPLICATION NUMBER: US/09/453,514A
CURRENT PILING DATE: 1999-12-01
FRIOR APPLICATION NUMBER: 09/144,611
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PatentIn version 3.0
SEQ ID NO 12
CURRENT FILING DATE: 2000-01-05
PRIOR APPLICATION NUMBER: 09/016,520
PRIOR FILING DATE: 1998-01-30
PRIOR FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
TYPE: DNA
TYPE: DNA
CORGANISM: Artificial Sequence
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; Sequence 12, Application US/09453514A
; Patent No. 6326199
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                      15; Conservative
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Best Local Similarity
Matches 15; Conserv
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LENGTH: 19
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Patent No. 6399754
GENERAL INFORMATION:
APPLICANT: Phillip Dan Cook
ITITLE OF INVENTION: Sugar Modified Oligonucleotides
NUMBER OF SEQUENCE: 37
CORRESPONDENCE ADDRESS:
ADDRESSEE Woodcock Washburn Kurtz Mackiewicz and No. 6399754ris
STREET: One Liberty Place - 46th Floor
CITY: Philadelphia
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                                                                                                                                                    Length 19;
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1.0%; Score 13.8; DB 1;
Best Local Similarity 88.2%; Pred, No. 1.3e+02;
Matches 15; Conservative 0; Mismatches 2;
                                                                                                                                                1.0%; Score 13.8; DB 1;
88.2%; Pred. No. 1.3e+02;
cive 0; Mismatches 2;
                                                           NAME/KEY: misc feature; OTHER INFORMATION: No. 6326199el Sequence US-09-453-514A-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ISIS-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDLUM TYPE: 3.5 inch disk, 720 Kb
COMPUTER: 1M PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/135,202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US/09/135,202
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/471,973
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-449-731-49/c
; Sequence 49, Application US/08449731
; Patent No. 6413727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEI/ Action 1 Total NAME: Joseph Lucci REGISTRATION NUMBER: 33,307 REFERENCE/DOCKET NUMBER: ISIS TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                    1141 GCCTTTTTTTTTTT 1157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1141 GCCTTTTTTTTTTT 1157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 33: SEQUENCE CHARACTERISTICS:
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19 GCGTTTTTTTTTTT
                                                                                                                                           Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     215-568-3439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             U.S.A.
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CURRENT FILING DATE: 1997-02-11
NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin version 3.1
SEQ ID NO 29
LENGTH: 19
                                                                                                       TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                   19 GCGTTTTTTTTTTT 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17 crcrccercrecceac
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 15; Conserval
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
Matches 15; Conserva
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                                                                                                                                                                                  NAKAMURA, YUSUKE
THILYGRIS, ANDREE
TILLE OF INVENTION: INDENE AND SONATIC MUTATIONS OF APC
GENE IN COLORECTAL CANCER IN HUMANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: RIOPRY disk
COMPUTER: IND PC compatible
COMPUTER: IND PC compatible
COMPUTER: DATEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/449,731
FILING DATE: 25-May-1995
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
RELEASE 24-May-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Cook, Phillip D.
APPLICANT: Monia, Brett
APPLICANT: Marin, Pierre
APPLICANT: Martin, Pierre
APPLICANT: Martin, Sayar-Modified Gapped Oligonucleotides
FILE OF INVENTION: SAyar-Modified Gapped Oligonucleotides
CURRENT APPLICATION NUMBER: US/08/802,331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
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Pred. No. 1.3e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 1107.46943
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9299
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 102
CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Allegretti, LTD
STREET: 1001 G Street, NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
SEQUENCE DESCRIPTION: SEQ ID NO: 49:
                                                                              GRODEN, JOANNA
HEDGE, PHILIP J.
JOSLYN, GEOFF
KINZLER, KENNETH
MARKHAM, ALEXANDER F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 29, Application US/08802331
Patent No. 6451991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100 ACAACCCCGGAGGCGCA 116
GENERAL INFORMATION:
APPLICANT: ALBERISEN, HANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17 ACAACCCAGGAGCCGCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 88.2%;
Matches 15; Conservative
                                                             CARLSON, MARY
                                                                                                                                                                                                                                                                                                                                             CITY: Washington STATE: D.C.
                                        ANAND,
                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORIGINAL SOURCE:
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US-08-802-331-29/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ·.
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                                                         1.0%; Score 13.8; DB 1; Length 19; larity 88.2%; Pred. No. 1.3e+02; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/375,318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Verna, James M.
REGISTRATION NUMBER: 33,287
REFRENCE/DOCKET NUMBER: 920010.571C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSE: SEED and BEERY LLP
STREET: 701 Fifth Ave, Suite 6300
CITY: Seattle
STATE: Washington
; OTHER INFORMATION: No. 6451991el Sequence
US-08-802-331-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 29:
                                                                                                                                                                                                                                    FILING DATE: 16-Aug-1999
CLASSIFICATION: <UNKNOWN>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        682-6031
                                                                                                                                       1141 GCCTTTTTTTTTTT 1157
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     581 CCCTCCGTCTGCCCCC 597
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STRANDEDNESS: single
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19 GCGTTTTTTTTTTTG
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 bases
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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US-08-832-021-49
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                                                                                                                                  APPLICANT: Tanzi, Rudolph E.
Schelberg, Gerard D.
Wasco, Wilma
Levy-Lahad, Ephrat
Bird, Thomas D.
Galas, David J.
TITLE OF INVENTION: CHRONSOME 1 GENE AND GENE PRODUCTS RELATED TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:

APPLICANT: Phillip Dan Cook
APPLICANT: A. Kawasaki.
TITLE CN T. A. Kawasaki.
TITLE CN F. FUNDALION: 2'-Modified Oligonucleotides
NUMBER OF SEQUENCES: 3'
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 6531584ris
STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Verna, James M.
REGISTRATION NUMBER: 33,287
REFERENCE/DOCKET NUMBER: 920010.571C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BEERY LLP
STREET: 701 Fifth Ave, Suite 6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 43:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 16-Aug-1999
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
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US-08-189-283-33/c
'Sequence 33, Application US/09389283
'Patent No. 6531584
                                                                         Sequence 43, Application US/09375318 Patent No. 6468791 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     581 CCCTCCGTCTGCCCCCC 597
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INFORMATION FOR SEQ ID NO: 43:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 88
                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 88.23
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             U.S.A.
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ZIP: 19103
                                   RESULT 133
US-09-375-318-43/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.0%; Score 13.8; DB 1;
88.2%; Pred. No. 1.3e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 76, Application US/09302681
Patent No. 6441149
GENERAL INFORMATION
APPLICANT: Herrnstadt, Corrina
APPLICANT: Ghosh, Soumitra S.
APPLICANT: Glevenger, William
APPLICANT: Clevenger, William
APPLICANT: Pahy, Eoin F.
TITLE OF INVENTION: DIAGNOSTIC METHOD BASED ON
TITLE OF INVENTION: QUANTIFICATION OF EXTRAMITOCHC
FILE REPERENCE: 660088.416C1
CURRENT APPLICATION NUMBER: US/09/302,681
CURRENT PILIG DATE: 1999-04-30
                                                                                                                                                                                                                APPLICALL...
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Joseph Lucci
REGISTRATION NUMBER: 33,307
REFERENCE/DOCKET NUMBER: ISIS-2004
TELECHONE: 215-568-3100
TELEFHONE: 215-568-3100
TELEFRAX: 215-568-3439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 108
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 76
LENGTH: 21
MEDIUM TYPE: 3.5 inch disk, 720 Kb COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WordPerfect 5.1 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/389,283
                                                                                                                                            FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/035,357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     268 TGGCTGATCAAAGAGGAAGC 287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 recercarreascactarec 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
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93.3%; Pred. No. 1.2e+02;
tive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILING DATE: 1997-11-13
PRIOR PILING DATE: 1997-11-24
PRIOR PILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 60/075945
PRIOR PILING DATE: 1998-02-25
PRIOR APPLICATION NUMBER: 60/078910
                                                                                                                                                                                                                  Sequence 493, Application US/09996243
Patent No. 6478825
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/065186
FILING DAFE: 1997-11-12
APPLICATION NUMBER: 60/065311
FILING DAFE: 1997-11-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/087106
FILING DATE: 1998-05-28
APPLICATION NUMBER: 60/087607
FILING DATE: 1998-06-02
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FILING DATE: 1998-06-04
APPLICATION NUMBER: 60/088025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/084600
FILING DATE: 1998-05-07
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FILING DATE: 1998-06-02
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FILING DATE: 1998-06-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Godowski, Paul J.
Grimaldi, J. Christopher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Watanabe, Colin K.
Williams, P. Mickey
Wood, William I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Roy, Margaret Ann
Stewart, Timothy A.
Tumas, Daniel
                                                                       237 GGCATCTGCATCTGG 251
                                                                                                                3 GGCATCTGCTTCTGG 17
                                                                                                                                                                                                                                                                                                                                                                                    Ferrara, Napoleone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gerritsen, Mary E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Paoni, Nicholas F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gurney, Austin L.
                                  14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kljavin,Ivar J.
Napier,Mary A.
                                                                                                                                                                                                                                                                                                                        Botstein, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Goddard, Audrey
                                                                                                                                                                                                                                                                                                                                             Desnoyers, Luc
Eaton, Dan L.
                                                                                                                                                                                                                                                                                                                                                                                                        Fong, Sherman
        Best Local Similarity
Matches 14; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pan, James
                                                                                                                                                                                                   US-09-996-243-493
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Patent No. 6043088
GENERAL INFORMATION:
APPLICANT: Bookstein, Robert
APPLICANT: Baccs, William B.
TITLE OF INVENTION: A No. 6043088el Prostate/Colon Tumor Suppressor
TITLE OF INVENTION: Gene Located on Human Chromosome 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
Sequence 49, Application US/08B32021
Patent No. 60459B
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Pardinas, J.
APPLICANT: Practy, S.
APPLICANT: Practy, S.
TILLE OF INVENTION: IMPROVED TECHNIQUE FOR DIFFERENTIAL DISPLAY
FILE REPERENCE: 1989-382
CURRENT APPLICATION NUMBER: US/08/832,021
CURRENT FILING DATE: 1999-04-02
NUMBER OF SEQ ID NOS: 64
SEQ ID NO 49
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Description of Artificial Sequence: primer US-08-832-021-49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 15,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/445,515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
CITY: San Diego
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.0%; Score 13.4; DB 1; 93.3%; Pred. No. 87;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFRENCE/DOCKET NUMBER: P-C.
TELECOMMINICATION INFORMATION:
TELEPAX: (619) 535-9901
TELEPAX: (619) 535-9949
INFORMATION FOR SEQ ID NO: 37:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1145 TITITICITITIGGA 1159
                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 5:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell a
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Best Local Similarity
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CLASSIFICATION:
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US-08-445-515-37
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APPLICANT: Wood, William I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same FILE REFERENCE: P2730P1C13
FILE REFERENCE: P2730P1C13
CURRENT APPLICATION NUMBER: US/09/996,243
CURRENT FILING DATE: 2001.11-14
PRIOR APPLICATION NUMBER: 60/049787
PRIOR APPLICATION NUMBER: 60/049787
PRIOR APPLICATION NUMBER: 60/062250
PRIOR APPLICATION NUMBER: 60/062250
Gaps
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PRIOR APPLICATION NUMBER: 6/008026
PRIOR APPLICATION NUMBER: 6/008028
PRIOR PELING DATE: 1939-06-04
PRIOR APPLICATION NUMBER: 6/008029
PRIOR PELING DATE: 1939-06-04
PRIOR PELING DATE: 1939-06-04
PRIOR PELING DATE: 1939-06-04
PRIOR PELING DATE: 1938-06-04
PRIOR PELING DATE: 1938-06-05
PRIOR PELING DATE: 1938-06-07
PRIOR APPLICATION NUMBER: 6/008625
PRIOR APPLICATION NUMBER: 6/008626
PRIOR APPLICATION NUMBER: 6/008626
PRIOR APPLICATION NUMBER: 6/008626
PRIOR APPLICATION NUMBER: 6/008626
PRIOR PELING DATE: 1938-06-10
PRIOR PELING DATE: 1938-06

### PRIOR APPLICATION NUMBER (0/09022
### PRIOR APPLICATION NUMBER (0/09024
### PRIOR APPLICATION NUMBER (0/09024
### PRIOR APPLICATION NUMBER (0/09034
### PRIOR PRIING DATE 1998-06-23
### PRIOR APPLICATION NUMBER (0/09069
### PRIOR APPLICATION NUMBER (0/091626
### PRIOR PRIING DATE 1998-06-23
### PRIOR PRIING DATE 1998-07-07
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Gaps

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Fatent No. 6008048
GENERAL INFORMATION:
JAPPLICANT: Brett P. Monia
APPLICANT: ex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF EGR-1 EXPRESSION
FILE REFERENCE: RTS-0028
CURRENT APPLICATION NUMBER: US/09/205,921A
NUMBER OF SEQ ID NOS: 47
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Brenda F. Baker
APPLICANT: Lex M. Cowsert
ITILE OF INVENTION: ANTISENSE MODULATION OF SENTRIN EXPRESSION
FILE REFERENCE: RE2-0026
                                                                                                                                                                                                                                                                                                                                                                          Length 18;
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                                                                                                                                                                                                                                                                                                                                                                    Query Match
1.0%; Score 13.4; DB 1;
Best Local Similarity 33.3%; Pred. No. 1.4e+02;
Matches 5; Conservative 9; Mismatches 1;
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1.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 1.4e+02;
Matches 14; Conservative 0; Mismatches 1;
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, OTHER INFORMATION: Antisense Oligonucleotide
US-09-213-768-47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/213,768
CURRENT FILING DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 47
SEQ ID NO 47
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US-09-213-768-47/c
Sequence 47, Application US/09213768
Patent No. 5985664
                TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: 67-3510
INFORMATION FOR SEQ ID NO: 2595:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDENNESS: single
TOPOLOGY: linear
US-08-585-684B-2595
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1111 GTTTTCTGTTTAATT 1125
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Best Local Similarity 93.3
Matches 14; Conservative
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APPLICANT: Structure, Jun
APPLICANT: Bacobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Layol, 2(37/198)
CURRENT APPLICATION NUMBER: US (9/005,974
PRIOR APPLICATI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2595, Application US/08585684B
Fatent No. 5877021
GENERAL INFORMATION:
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
NUMBER OF SEQUENCES: 2751
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Indels
                                                                Sequence 5187, Application US/09371772B Patent No. 6566127 GENBRAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FasteSO Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,684B
FILING DATE: January 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                218/078
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STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1154 TTTGGAAGTAAAGCA 1168
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/000
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
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US-09-371-772B-5187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Los Angeles
STATE: California
                                                                                                                                                                                               Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 10; Conserv
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      RESULT 139
US-09-371-772B-5187
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                                                                                                                                                                                               APPLICANT:
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                 US-09-038-073-2595
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                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF G-ALPHA-16 EXPRESSION
FILE REPERENCE: FTS-003
CURRENT APPLICATION NUMBER: US/09/205,143
CURRENT FILING DATE: 1998-12-03
NUMBER OF SEQ ID NOS: 87
SEQ ID NO 11
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 1.0%; Score 13.4; DB 1; Length 18; Best Local Similarity 93.3%; Pred. No. 1.46+02; Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
NUMBER OF SEQUENCES: 2751
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Sal West Fifth Street
STREET: Los Angeles
STREET: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Antisense Oligonucleotide US-09-205-143-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FRACESQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US/09/038,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REPRENCE/DOCKET NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 2595, Application US/09038073; Patent No. 6194150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/038, CRILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/585,684
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                 ; Sequence 11, Application US/09205143; Patent No. 6107091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1336 GIGTITCAGGCAGGG 1350
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(213) 955-0440
348 CAGTGGCCGAGTGAG 362
                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16 GTGTTTCAGGCAAGG 2
                                           15 CAGTGGCCTAGTGAG 1
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                                                                                                        RESULT 143
US-09-205-143-11/c
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US-09-038-073-2595
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1.0%; Score 13.4; DB 1; Length 18;
Best Local Similarity 93.3%; Pred. No. 1.4e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                        1.0%; Score 13.4; DB 1; Length 18; 13.3%; Pred. No. 1.4e+02; ve 9; Mismatches 1; Indels
                                                                                                                                                                                                                                                 US-09-632-580A-34

US-09-632-580A-34

Sequence 34, Application US/09632580A

PREAR NO. 625511

GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: LEX M. COWSET

TITLE REPRENCE:
CURRENT APPLICATION NUMBER: US/09/632,580A

CURRENT FILING DATE: 2000-07-31

NUMBER OF SEQ ID NOS: 93

SEQ ID NO 34

LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ) OTHER INFORMATION: Antisense Oligonucleotide US-09-632-580A-34
                                                                                                                                                                                  1111 GITTICIGITIAATT 1125
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             552 GGCAGGCATGCACAC 566
                                                                                                                                         33.3%;
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1 GUUTUCUGUCUAAUU 15
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANBEDNES: single
TOPOLOGY: linear
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us09904568-3.rni

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Query Match
Best Local Similarity
Matches 14; Conserva
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          US-08-053-131-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: 8
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Patent No. 5661016
GENERAL INFORMATION:
APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5661016-Human Animals for TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 197
                                                                                                                                                                                                                                                                                                                 Length 19;
                                                                                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew STREET: One Market Plaza, Steuart Tower, Suite 200 CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Version #1.25
                                                                                                                                                                                                                                                                                                               1.0%; Score 13.4; DB 1; 93.3%; Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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ZUP: 94105

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: LATION DATA:
APPLICATION NUMBER: US/08/053,131
FILING DATE: 26-APR-1993
CLASSIPICATION DATA:
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/810,279
FILING DATE: 118-MR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 14643-9-3
TELECOMMUNICATION NUMBER: 14643-9-3
TELECOMMUNICATION NUMBER: 14643-9-3
TELECOMMUNICATION NUMBER: 14643-9-3
TELEPHONE: 415-356-2400
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 14643-5
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-543-9600
TELEPHONE: 415-543-9600
INFORMATION FOR SEQ ID NO: 860ENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                391 GTGGCAGCAATGGCC 405
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                        14; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
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US-08-053-131-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
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                                            Gaps
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Sequence 16, Application US/08645641

PARENT NO. 5719032

GENERAL INFORMATION:

APPLICANT: Lonberg, Nils

APPLICANT: ANY ROBERT M.

TITLE OF INVENTION: Transgenic No. 5719032-Human Animals for TITLE OF INVENTION: Producing Heterologous Antibodies

NUMBER OF SEQUENCES: 150

CORRESPONDENCE ADDRESS:

ADDRESSEE: William M. Smith

STREET: Two Embarcadero Center, 8th Floor

CITY: San Francisco
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APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5789650-Human Animals for
1.0%; Score 13.4; DB 1; Length 19; 93.3%; Pred. No. 1.5e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA

ZIP: 94111-3834

ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/645,641

FILING ATTE: 20-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 13.4; DB 1;
Pred. No. 1.5e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 20-MAY-1996
CLASSIFICATION: 800
PRICA APPLICATION DATA:
APPLICATION NUMBER: US 07/904,068
FILING DATE: 3-UN-1992
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 31,244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 16, Application US/07853408B Patent No. 5789650 GENERAL INFORMATION:
                                                                                  391 GTGGCAGCAATGGCC 405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: DNA (primer)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 93.3%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     391 GTGGCAGCAATGGCC 405
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                                          14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  415-326-2422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19 base pairs
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Patent No. 5874299
| Fatent No. 5874299 | GENERAL INFORMATION:
| FATENT LINE OF INVENTION: | FAPPLICANT: Lonberg, Nils | APPLICANT: Kay, Robert M. | TITLE OF INVENTION: | Producing Heterologous Antibodies | TITLE OF INVENTION: | Producing Heterologous Antibodies | NUMBER OF SEQUENCES: | ADDRESSEE: | Milliam M. Smith | STREET: | One Market Plaza, Steuart Tower, Suite 2000 | STREET: | California | STATE: | California | Cal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/800,353
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
1.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 1.5e+02;
Matches 14; Conservative 0; Mismatches 1;
                                                                                                                         PRIOR APPLICATION DAMBER: US
PELING DATE: 23-UN-1992
PRIOR APPLICATION DATE:
PRIOR APPLICATION DATE:
APPLICATION NUMBER: US
FILING DATE: 18-MAR-1992
PRIOR APPLICATION DATE:
APPLICATION NUMBER: US 07/810,279
FILING DATE: 17-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 14643-9-4
TELEPHONE: 415-326-240
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
MUMBER: MUMBER: 1000 MUMBER: 
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/834,539
FILING DATE: 1992-02-05
ATTORNEY/AGENT INFORMATION:
         APPLICATION NUMBER: US 07/990,860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Smith, William M.
REGISTRATION UNDRER: 30,223
REFERENCE, DOCKET NUMBER: 1464:
TELECOMMUNICATION INFORMATION:
TELEFONE: 415-543-9600
TELEFAX: 415-543-5043
INFORMATION FOR SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   , MOLECULE TYPE: DNA (primer) US-08-096-762-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             391 GTGGCAGCAATGGCC 405
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                                                          16-DEC-1992
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STRANDEDNESS: single
TOPOLOGY: linear
                                                     FILING DATE: 16-DEC-1
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U
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APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5814318-Human Animals for
TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 210
CORRESPONDENCE ADDRESS:
ADDRESSEE: Towneend and Townsend Khourie and Crew
STREET: One Market Plaza, Steuart Tower, Suite 200
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.0%; Score 13.4; DB 1; Length 19; 93.3%; Pred. No. 1.5e+02; Live 0; Mismatches 1; Indels
TITLE OF INVENTION: Producing Heterologous Antibodies NUMBER OF SEQUENCES: 150
CORRESPONDENCES: ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION DATA:

APPLICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: US/07/853,408B

FILING DATE: 19920318

CLASSIFICATION: 800

ATTORNEY/AGENT INFORMATION:

NAME: Smith, William M.

REGISTRATION NUMBER: 30,223

REGISTRATION NUMBER: 30,223

REGISTRATION NUMBER: 36,223

REGISTRATION NUMBER: 31,223

RELEPHONE: 415-326-2400

TELEFHONE: 415-326-2400

TELEFACOMUNICATION: 16: SEQUENCE CHARACTERISTICS:

LINFORMATION FOR SEQ ID NO: 16: SEQUENCE CHARACTERISTICS:

LINFORMATION FOR SEQ ID NO: 16: SEQUENCE CHARACTERISTICS:

TYPE: NUCLEIC ACID

STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/096,762
FILING DATE: 22-JUL-1993
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                            ZIP: 94105
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         391 GTGGCAGCAATGGCC 405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear; MOLECULE TYPE: DNA (primer)
US-07-853-408B-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 GIGGCCGCAAIGGCC 18
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Best Local Similarity 93.33
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                               USA
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US-08-096-762-16
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FILING DATE: 02-DEC-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/HSG7/77
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 09-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 10-OCT-1
PRIOR APPLICATION NUMBER: 1
                                                                                                                                                                                                                                               CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                     US-09-042-353-184
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Patent No. 5877397
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Transgenic No. 5877397-Human Animals for TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 150
CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%; Score 13.4; DB 1; Length 19; 93.3%; Pred. No. 1.5e+02;
                                                                                                                                                   Length 19;
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                                                                                                                                                                                       1; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURSENI APPLICATION DATA:
APPLICATION NUMBER: US/08/308,865
FILING DATE:
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/145,707
FILING DATE: 23-JUN-1992
ATORNEY/AGENT INFORMATION:
REFILENCE SMITH, WIlliam M.
REFERENCE/COKET NUMBER: 30,223
REFERENCE/COKET NUMBER: 14643-9-1-1
TELEPHONE: A15-356-2400
                                                                                                                                                 Score 13.4; DB 1;
Pred. No. 1.5e+02;
                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                       TOPOLOGY: linear

MOLECULE TYPE: DNA (genomic)
US-08-800-353-8
                                                                                                                                                   1.0%;
                                                                                                                                                                                                                           391 GTGGCAGCAATGGCC 405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; MOLECULE TYPE: DNA (primer)
US-08-308-865-16
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
               1: 19 base pairs
nucleic acid
DEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best_Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                     14; Conservative
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: One Main.C.
CITY: San Francisco
STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                 TYPE: nucleic
STRANDEDNESS:
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US-08-308-865-16
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GENERAL INFORMATION:

APPLICANT: Lonberg, Nils
APPLICANT: Robert M.
TITLE OF INVENTION: Transgenic No. 6255458-Human Animals for TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 421
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LIP
STREET: Two Embarcadero Center, Bighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/042,353
FILING DATE: 13-MAR-1998
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/810,279
FILING DATE: 17-DEC-1991
PRIOR APPLICATION NUMBER: US 07/853,408
FILING DATE: 17-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/953,408
FILING DATE: 18-MAR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/990,860
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10-OCT-1996
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FILING DATE: 07-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/544,404
FILING DATE: 10-CCT-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/728,463
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/096,762
FILING DATE: 22-UUL-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/155,301
FILING DATE: 18-NOV-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/155,301
FILING DATE: 03-DEC-1993
PRIOR APPLICATION DATA:
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FILING DATE: 10-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION THE PRIOR THE PRIO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94111.3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
. Sequence 184, Application US/09042353; Patent No. 6255458
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WO PCT/US97/21803

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Gaps
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GENERAL INFORMATION:

APPLICANT: Lonberg, Nils

APPLICANT: Ray, Robert M.

TITLE OF INVENTION: Transgenic Non-Human Animals Capable of

TITLE OF INVENTION: Producing Heterologous Antibodies

NUMBER OF SEQUENCES: 75

CORRESPONDENCE ADDRESS:

ADDRESSEE: William M. Smith

STREET: One Market Plaza, Steuart Tower, Suite 2000

CITY: San Francisco

STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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1.0%; Score 13.4; DB 1; Length 19;
Best Local Similarity 93.3%; Pred. No. 1.5e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           1.0%; Score 13.4; DB 1; Length 19; 93.3%; Pred. No. 1.5e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Lan Popoff
APPLICANT: Lan Popoff
APPLICANT: Lax M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF PARP EXPRESSION
FILE REFERENCE: RTS-0150
CURRENT APPLICATION NUMBER: US/09/517,467B
CURRENT FILING DATE: 2001-03-02
PRIOR FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 345
LENGTH: 19
                               NAME: Serafini, Andrew T.
REGISTRATION NUMBER: 41,303
REFERENCE/DOCKET NUMBER: 014643-009030US
                                                                                                                                                                                                                                                                                  TOPOLOGY: linear TOPOLOGY: linear is MOLECULE TYPE: DNA SEQUENCE DESCRIPTION: SEQ ID NO: 32: US-08-758-417A-32
                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHOX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 9, Application US/09517467B
; Patent No. 6451602
; GENERAL INFORMATION:
         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        391 GTGGCAGCAATGGCC 405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: PCR Primer
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Matches 14; Conservative
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Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 6300129-Human Animals for
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ò
                                                                                                                                                                                                                                                                                                                                                                                                      1.0%; Score 13.4; DB 1; Length 19; 93.3%; Pred. No. 1.5e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Producing Heterologous Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA

COMPUTER READABLE FORM.

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
FILING DATE: 01-DEC-1997
ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REGISTRANCE/DOCKET NUMBER: 014643-009040US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 184:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 0.2-bec-1996
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA: 0.8/728,463
FILING DATE: 10-OCT-1996
APPLICATION NUMBER: US 08/544,404
FILING DATE: 10-OCT-1995
APPLICATION NUMBER: US 08/542,322
APPLICATION NUMBER: US 08/52,322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PILING DATE: 10-DEC-1993
APPLICATION NUMBER: US 08/161,739
FILING DATE: 03-DEC-1993
APPLICATION NUMBER: US 08/155,301
FILING DATE: 18-NOV-1993
APPLICATION NUMBER: US 08/096,762
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APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 07-DEC-1994

PELLING DATE: 09-MAR-1994

FILING DATE: 09-MAR-1994

APPLICATION NUMBER: US 08/165,699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 32, Application US/08758417A; Patent No. 6300129; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 391 GTGGCAGCAATGGCC 405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Greececharescc 18
                                                                                                                                                                                                                               LENGTH: 19 base pairs

TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 14; Conserv?
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 154
US-08-758-417A-32
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Gaps

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GENERAL INFORMATION:
APPLICANT: IVETSEN, PATRICK L.
TITLE OF INVENTION: Methods and Compositions for Their
TITLE OF INVENTION: Methods and Compositions for Their
TITLE OF INVENTION: Use to Treat Metal Toxicity
                                                                                Score 13.4; DB 1; Length 19; Pred. No. 1.5e+02; 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA

ZIP: 23187-3609

COMPUTER READBLE FORM:
MEDIUM TYRE: Diskette, 5.25 inch, 360 Kb storage COMPUTER: IBM-compatible 486/33

OPERATING SYSTEM: Mb.DOS 5.0

SOFTWARE: IBM-compatible 486/33

OPERATING SYSTEM: McdPerfect 5.1

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/759,841C

FILING DATE: 12 September 1991

CLASSIFICATION: 514

PRIOR APPLICATION DATA: none
ATTORNEY/AGENT INFORMATION:
NAME: FLOYd, John P.
REGISTRATION NUMBER: 19,528

REGISTRATION NUMBER: 19,528

REGISTRATION NUMBER: 63031

TELECOMMUNICATION INFORMATION:
TELEPHONE: (804) 220-0930
INFORMATION INFORMATION:
TELEPHONE: (804) 220-0930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PUBLICATION INFORMATION:
AUTHORS: Karin, M., and Richards, R. I.
TITLE: Human metallothionein genes-primary
TITLE: structure of the Metallothionein-II
TITLE: a related processed gene
JOURNAL: Nature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ANTI-SENSE: yes
POSITION IN GENOME:
CHROMOSOME/SEGMENT: 16q21-q22.1.
MAP POSITION: 844 through 861
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2, Application US/07759841C Patent No. 5618796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSER: John P. Floyd, Esq.
STREET: P.O. Box 3609
CITY: Williamsburg
STATE: Virginia
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EDNESS: single stranded
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MOLECULE TYPE: CDNA to a mRNA
HYPOTHETICAL: no
                                                                                   1.0%;
                                                                                                                                                                                                         391 GIGGCAGCAAIGGCC 405
                                                                                                                                                                                                                                                                         4 Gracececharedec 18
                                                                      Query Match
Best Local Similarity 93.3?
Matches 14; Conservative
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Best Local Similarity 83.3
Matches 15; Conservative
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                    PCT-US92-10983-16
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ISSUE: 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic Non-Human Animals for TITLE OF INVENTION: Producing Heterologous Antibodies; NUMBER OF SEQUENCES: 152
CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
STREET: California
COUNTRY: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: THEN PC compatible
OPERATING SYSTEM: PC-DOS/WS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/06185
FILING DATE: 19910828
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 87654
REFERENCE/DOCKET NUMBER: 14643-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM;

COMPUTER READABLE FORM;

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US92/10983

FILING DATE: 19921217

ATTORNEY AGENT INFORMATION:

NAME: Smith, William M.

REGISTRATION NUMBER: 14643-9-2

REFERENCE/DOCKET NUMBER: 14643-9-2

TELEPHONE: 415-326-2400

TELEFRA: 415-326-2400

TELEFRA: 415-326-2402

INFORMATION FOR SEQ ID NO: 16: SEQUENCE CHARACTERISTICS:

LENGTH: 19 base pairs

TYPE: NUCLEIC ACID

STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-543-9600
TELEPAX: 415-543-9603
INFORMATION FOR SEQ ID NO: 8: SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             391 GTGGCAGCAATGGCC 405
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Sequence 7, Application US/09073465
Fatent No. 605478
Fatent Nober Nober
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                                US-09-339-964-33
; Sequence 33, Application US/09339964
; Patent No. 6025198
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          869 TCCCCACAGCCAAGTTCC 886
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
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RESULT 160
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                                                                                                                                                                                                                       Sequence 3, Application US/07759841C

Patent No. 5518796

GENERAL INFORMATION:

TITLE OF INVENTION: No. 5618796el Metal Binding Agents, and TITLE OF INVENTION: Methods and Compositions for Their TITLE OF INVENTION: Use to Treat Metal Toxicity

NUMBER OF SEQUENCES: 3

CORRESPONDENCE ADDRESS:

ADDRESSEE: John P. Ployd, Esq.

STREET: P.O. Box 3609
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 1.5e+02; ive 0; Mismatches 3; Indels
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POSITION GENOME:
CHROMOSOME/SENSENT: 16q21-q22.1.
CHROMOSOME/SENSENT: 16q21-q22.1.
MAP POSITION: 844 through 861
UNITS:
PUBLICATION: M., and Richards, R. I.
TITLE: Human metallothionein genes-primary
TITLE: structure of the Metallothionein-II gene and
TITLE: a related processed gene
VOLUME: 299
ISSUE: 29
ISSUE: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 5.25 inch, 360 Kb storage COMPUTER: IBM-compatible 486/33
OPERATING SYSTEM: MS-DOS 5.0
SOFTWARR: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/759,841C
FILING DATE: 12 September 1991
CLASSIFICATION: 5.4
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION OF SEPTEMBER: US/07/759,841C
FILING DATE: 12 September 1991
CLASSIFICATION DATA: none
ATTORNEY/AGENT INFORMATION:
NAME: Floyd John P.
REGISTRATION UNDERF: 19,528
                                                                                                                                                                                                                                                                                                                                                                                                                                                         J. Leat.
J. L. L. D. Dox 3609
CITY: Williamsburg
STATE: Virginia
COUNTRY: Ush
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               63031
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REFRENCE DOCKET UNDRER: 6303:
TELECOMMUNICATION INFORMATION:
TELEPAX: (804) 220-0930
TELEPAX: (804) 220-0930
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 nucleotide bases
TYPE: nucleic acid
STRANDENESS: single
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525 GCCGGAGGAGCAGCTGGG 542
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                                                          1 GGCGCAGGAGCAGTTGGG 18
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                RESULT 159
US-07-759-841C-3/c
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US-07-759-841C-3
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Patent No. 6040179

GENERAL INFORMATION

TITLE OF INVENTION: ANTISENSE MODULATION OF G-ALPHA-I2 EXPRESSION
FILE REPERENCE: RTS-0064

CURRENT APPLICATION WHERE: US/09/339,993A

CURRENT FILING DATE: 1999-06-25

NUMBER OF SEQ ID NOS: 47

LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ..
APPLICANT: C. Frank Bennett
APPLICANT: C. Frank Bennett
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF SHIP-2 EXPRESSION
FILE REFERENCE: RTS-0065
CURRENT APPLICATION NUMBER: US/09/339,964
CURRENT FILING DATE: 1999-06-25
NUMBER OF SEQ ID NOS: 47
SEQ ID NO 33
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
1.0%; Score 13.2; DB 1; Length 18;
Best Local Similarity 83.3%; Pred. No. 1.5e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Antisense Oligonucleotide US-09-339-964-33
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us09904568-3.rni

Page 59

LENGIH: 18

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STRANDEDNESS:
           RESULT 165
US-08-795-430-31
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US-09-487-444-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: C. Frank Bennett
APPLICANT: Brett P. Monia
APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTIENSE MODULATION OF NF-KAPPA-B P65 SUBUNIT EXPRESSION
FILE REFERENCE: RTS-0025
CURRENT APPLICATION NUMBER: US/09/199,859
CURRENT FILING DATE: 1998-11-25
SEQ ID NO 14
LENGTH: 18
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Sequence 31, Application US/09339775

Pattent No. 6063626

GENERAL INFORMATION:
APPLICANT: Lex M. Cowsert

TITLE OF INVENTION: ANTISENSE MODULATION OF G-ALPHA-I3 EXPRESSION
FILE REFERENCE: RTS-0069

CURRENT APPLICATION UNBER: US/09/339,775

CURRENT FILING DATE: 1999-06-24

NUMBER OF SEQ ID NOS: 47

LENGTH: 18
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0
                                                       ; OTHER INFORMATION: Description of Unknown Organism: Bacterial US-09-073-465-7
                                                                                                                1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 1.5e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 1.5e+02; Live 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.0%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 1.5e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Antisense Oligonucleotide US-09-339-775-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 14, Application US/09199859
; Patent No. 6069008
; GENERAL INFORMATION:
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                                                                                                                                                                                       882 GTTCCAGGAGCTGCGGTA 899
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
TYPE: DNA
ORGANISM: Unknown Organism
                                                                                                                                                        Conservative
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Best Local Similarity
Thes 15; Conserva
                                                                                                      Query Match
Best Local Similarity
Matches 15; Conserv
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Best Local Similarity
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US-09-199-859-14
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                                     FEATURE:
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Sequence 31, Application US/08795430
| Patent No. 6130071
| GENERAL INFORMATION:
| APPLICANT: Alitalo, Kari
| APPLICANT: Joukov, Vladimir
| TITLE OF INVENTION: Vascular Endothelial Growth Factor C (VEGF-C)
| TITLE OF INVENTION: Protein and Gene, Mutants Thereof, and Uses Thereof
| NUMBER OF SEQUENCES: 57
| CORRESPONDENCE ADDRESS: ADDRESS: ADDRESSE: Marshall, O'Toole, Gerstein, Murray & Borun
| STREET: 6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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                                                                                                                                                                                                                                Query Match
1.0%; Score 13.2; DB 1;
Best Local Similarity 83.3%; Pred. No. 1.5e+02;
Matches 15; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
REICR APPLICATION DATA:
APPLICATION NUMBER: PCT/F196/00427
FILING DATE: 01-AUG-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/671,573
FILING DATE: 28-JUN-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/601,132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 38,153
REFERENCE/DOCKET NUMBER: 28967/33691
TELECOMMUNICATION INFORMATION:
TRIEDENNE: 312/474.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 14-FEB-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/58,895
FILING DATE: 12-GAN-1996
FRICH APPLICATION NUMBER: 08/510,133
FILING DATE: 01-AUG-1995
PRIOR APPLICATION NUMBER: 08/340,011
FILING DATE: 14-NOV-1994
ATTORNEY/AGENT INFORMATION:
NAME: GASS, DAVIG A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                380 Trccrcagaggragcag 397
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 25-3856
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               312/474-0448
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EDNESS: single
                                                                                                                                                                                                                        CITY: Chicago
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Page 60

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NAME/KEY: misc_feature

// LOCATION: 1..18

// OTHER INNORMATION: upstream amplification primer for SEQ 218, SEQ 295, SEQ 219, US-09-218-207-354
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| Patent No. 6346398
| GENERAL INFORMATION |
| APPLICANT: Pavco, Pamela |
| APPLICANT: Bacched, Dan T. |
| APPLICANT: Stinchcomb, Dan T. |
| TITLE OF INVENTION: TREATMENT OF DISEASES OR |
| TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS |
| TITLE OF INVENTION: GROWTH FACTOR |
| NUMBER OF SEQUENCES: 8502 |
| CORRESPONDENCE ADDRESS: ADDRESS: ADDRESSE: LYON & LYON |
| STREET: SJA West Fifth Street |
| SJA West Fifth Stree
                                   APPLICANT: Ilya, Chumakov
APPLICANT: Bougueleret, Lydie
TITE OF INVENTION: Prostate cancer gene
FILE REFERENCE: GENSET.018CP1
CURRENT APPLICATION NUMBER: US/09/218,207
CURRENT FILING DATE: 1998-12-22
EARLIER APPLICATION NUMBER: 08/996,306
EARLIER FILING DATE: 1997-12-22
EARLIER FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 578
SOFTWARE: Patent.pm
SOFTWARE: Patent.pm
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORL PEFFECT 5.1
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APPLICATION NUMBER: US/08/584,040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60/005,974
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: January 11, 1996
CLASSIFICATION: 514
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          APPLICANT: Blumenfeld, Marta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 83.3
Matches 15; Conservative
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Homo Sapiens
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STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 XY: U.S.A. 90071-2066
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1 LOCATION: 1..18

2 OTHER INFORMATION: upstream amplification primer for SEQ 218, SEQ 295, SEQ 219, SEQ US-09-338-907-354
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                                                                                           APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF SMAD7 EXPRESSION
FILE REPERENCE: TTS-0133
CURRENT APPLICATION NUMBER: US/09/487,444
CURRENT FILING DATE: 2000-01-19
NUMBER OF SEQ ID NOS: 49
ESCO ID NO ID NOS: 49
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Antisense Oligonucleotide US-09-487-444-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Cohen, Daniel
APPLICANT: Cohen, Daniel
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Lydie
TITLE OF INVENTION: PROSTATE CANCER GENE
TITLE OF INVENTION: PROSTATE
CURRENT APPLICATION NUMBER: US/09/338,907
CURRENT FILING DATE: 1999-06-23
BARLIER APPLICATION NUMBER: 06/099,658
BARLIER PILING DATE: 1998-12-22
BARLIER PILING DATE: 1998-12-22
MUMBER OF SEQ ID NOS: 578
SEQ ID NO 354
LENGTH: 18
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Sequence 354, Application US/09218207
Patent No. 6346381
GENERAL INFORMATION:
APPLICANT: Cohen, Daniel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 354, Application US/09338907 Patent No. 6265546
Sequence 10, Application US/09487444
Patent No. 6159697
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               869 TCCCCACAGCCAAGTTCC 886
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 crccedcrecccaccc 18
                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 15; Conserva'
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US-09-338-907-354
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Best Local Similarity
                  Matches
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                                                                                                                                                                                                     Length 18;
                                                                                                                                                                                                                                     3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL GROWTH FACTOR
                                                                                                                                                                                                 1.0%; Score 13.2; DB 1; 72.2%; Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stochedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
                                                                                                                                                                                                                                     2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: BEAN Compatible
COMPATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 0/1005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 170
US-08-584-040-4454
; Sequence 4454, Application US/08584040
; Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                         142 CCGCTCGGCTCCGCTCCG 159
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2983:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                           1 ccucucagcuccucccca 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (213/ ...
TELEPHONE: (213/ 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                 | LENGTH: B base pairs | TYPE: nucleic acid | STRANDEDNESS: single | JOPOLGGY: linear | US-08-584-040-2983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Los Angeles
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                 Query Match
Best Local Similarity
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1.0%; Score 13.2; DB 1; Length 18;

Query Match

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                                   Indels
                                                                                                                                                                                                                                          us-u8-584-040-8393
; Sequence 8393, Application US/08584040
; Patent No. 6346389
; GENERAL INFORMATION:
    APPLICANT: Pavco, Pamela
    APPLICANT: McSwiggen, James
    APPLICANT: Stinchcomb, Dan T.
    APPLICANT: McSwiggen, James
    APPLICANT: Stinchcomb, Dan T.
    APPLICANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 13.2; DB 1;
Pred. No. 1.5e+02;
4; Mismatches 3;
61.1%; Pred. No. 1.5e+02;
tive 4; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           218/064
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US-09-355-700-31
; Sequence 31, Application US/09355700
; Patent No. 636L946
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/0
TELECOMMUNICATION INFORMATION:
TELERAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 8393:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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                                                                                                               1239 GCTGGACGTGGCCATGTG 1256
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Best Local Similarity 61.1%;
Matches 11; Conservative
                                   11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18 base pairs
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Los Angeles
California
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                                                                                                                                                                                                                                                                                                      RESULT 171
US-08-584-040-8393
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OTHER INFORMATION: Description of Artificial Sequence: No. 6451968el Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 1.0%; Score 13.2; DB 1; Length 18; Best Local Similarity 83.3%; Pred. No. 1.5e+02; Matches 15; Conservative 0; Mismatches 3; Indels
            APPLICANT: Monia, Brett P.
APPLICANT: Xu, Xiaoxing S.
TITLE OF INVENTION: ANTISENSE MODULATION OF TRAF EXPRESSION
FILE REFERENCE: ISPH-0321
CURRENT APPLICATION NUMBER: US/09/167,109
CURRENT FILING DATE: 1998-10-06
NUMBER OF SEQ ID NOS: 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.2; DB 1;
Pred. No. 1.5e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Curi, Peter Buchardt, Ole Ducholm, Kim D. Christensen, Leif Christensen, Leif WYFNTION: Linked Peptide Nucleic Acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANY: Christensen, Leif
TITLE OF INVENTION: Linked Peptide Nucleic of File Reference: ISIS1577
CURRENT APPLICATION NUMBER: US/08/275,951
CURRENT FILING DATE: 1994-07-15
READR APPLICATION NUMBER: 08/108,591
READR APPLICATION NUMBER: 08/088,658
READR APPLICATION NUMBER: 08/088,651
READR FILING DATE: 1993-07-02
READR APPLICATION NUMBER: 08/088,661
PRIOR FILING DATE: 1993-07-02
READR APPLICATION NUMBER: PCT/EP92/01219
READR APPLICATION NUMBER: 986/91
READR FILING DATE: 1991-05-22
READR APPLICATION NUMBER: 986/91
READR FILING DATE: 1991-05-22
READR APPLICATION NUMBER: 987/91
READR FILING DATE: 1991-05-24
READR APPLICATION NUMBER: 510/92
READR APPLICATION NUMBER: 510/92
READR APPLICATION NUMBER: 510/92
READR FILING DATE: 1991-05-24
READR FILING DATE: 1991-05-24
READR FILING DATE: 1991-05-24
READR RELING DATE: 1991-05-24
                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: antisense sequence US-09-167-109-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 33, Application US/08275951
Patent No. 6451968
GENERAL INFORMATION;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       471 GCAGGGGAAGCTGCCG 488
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                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Griffin, Michael
Coull, James M.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 83.33
Matches 15, Conservative
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Kiely, John
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                                                                                                                                                                                                                  SEQ ID NO 8
LENGTH: 18
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                         FEATURE:
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                        APPLICANT: Ludwig Institute for Cancer Research
Helsinki University Licensing
Alitalo, Kari(U.S. only)
Joukov, Vladimir (U.S. only)
TITLE OF INVENTION: Vascular Endothelial Growth Factor C (VEGF-C)
Protein and Gene, Mutants Thereof, and Uses Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                      ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & Borun STREET: 6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ·,
                                                                                                                                                                                                                                                                                                                               CITY: Chicago
STATE: Lilianois
COUNTRY: United States of America
ZIP: 60606-6402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/355,700
FILING DATE: 05-No. 6361946-1999
RILIA APPLICATION NUMBER: 08/795,430
FILING DATE: 05-FEB-1997
PRIDATE APPLICATION NUMBER: PCT/F196/00427
FILING DATE: 01-NG-1996
APPLICATION NUMBER: 08/601,132
FILING DATE: 14-FEB-1996
APPLICATION NUMBER: 08/601,132
FILING DATE: 12-JAN-1996
APPLICATION NUMBER: 08/601,133
FILING DATE: 12-JAN-1996
APPLICATION NUMBER: 08/601,133
FILING DATE: 12-JAN-1996
APPLICATION NUMBER: 08/510,133
FILING DATE: 01-AUG-1995
APPLICATION NUMBER: 08/340,011
FILING DATE: 14-NOV-1994
ATTORNEY/AGENT INFORMATION:
NAME: Cases David A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 1.0%; Score 13.2; DB 1; Length 18; Best Local Similarity 83.3%; Pred. No. 1.5e+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Gass, David A.
REGISTRATION NUMBER: 38,153
REFERENCE/DOCKET UNPBER: 28967/34140
TELECOMUNICATION INFORMATION:
TELEPHONE: 312/474-6300
TELEFAX: 312/474-6408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) SEQUENCE DESCRIPTION: SEQ ID NO: 31:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 8, Application US/09167109
Patent No. 6399297
GENERAL INFORMATION:
APPLICANT: Baker, Brenda F.
APPLICANT: Cowsert, Lex M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         380 TICCICCAGAGGIGGCAG 397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 TTCTTCCAAAGGTGTCAG 18
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INFORMATION FOR SEQ ID NO: 31:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                      NUMBER OF SEQUENCES:
GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-355-700-31
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APPLICANT: McGwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobed, Jaine
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFREENCE: MBHBOO, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1999-08-10
PRIOR PILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SEQ ID NO 4049
LENGTH: 18
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61.1%; Pred. No. 1.5e+02;
tive 4; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 13.2; DB 1;
Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4; Mismatches
                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 4049, Application US/09371772B ; Patent No. 6566127
                                                                                                                                                                                                       Sequence 2167, Application US/09371772B Patent No. 6566127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1239 GCTGGACGTGGCCATGTG 1256
                                142 CCGCTCGCTCCGCTCCG 159
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                                                                               1 cenebesecucences 18
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Best Local Similarity 61.1%;
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                     Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pavco, Pam
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Best Local Similarity
Matches 11; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-371-772B-2167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 178
US-09-371-772B-4049
                                                                                                                                                                                    US-09-371-772B-2167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: RNA
                                                                       셤
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Sequence 1411, Application US/09371772B

Patent NO. 6566127

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Baccow of Jame

APPLICANT: Baccobedo, Jaime

APPLICANT: Stinchcomb, Dan

APPLICANT: Scionchcomb, Dan

APPLICANT: Baccobedo, Jaime

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION NUMBER: US/09/371,772B

CURRENT APPLICATION NUMBER: US 60/005,974

PRIOR FILING DATE: 1995-01-08

NUMBER OF SEQ ID NOS: 14225

SOFUTANE: PatentIn version 3.0

SEQ ID NO 1411

LENGTH: 18
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                                                                                                                                                                    Sequence 6039, Application US/09422978

Patent No. 6537751

GENERAL INFORMATION:
APPLICANT: Blumenfeld, Marta
APPLICANT: Chunakov, Ilva
FILE REPERENCE: GENBET.020CP1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT FILING DATE: 1999-04-21
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER PILING DATE: 1998-11-23
EARLIER PILING DATE: 1998-14-21
EARLIER PILING DATE: 1998-14-21
EARLIER FILING DATE: 1998-14-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ) NAME/KEY: primer_bind

: LOCATION: 1..18

: OTHER INFORMATION: upstream amplification primer 99-8571 for SEQ 2105,

US-09-422-978-6039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ch 1.0%; Score 13.2; DB 1; Length 18; 1 Similarity 83.3%; Pred. No. 1.5e+02; 15; Conservative 0; Mismatches 3; Indels
1138 TATGCCTTTTTTTTT 1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        73
                                              1 Trircrirititiciti
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 72.2
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: Homo sapiens
US-09-371-772B-1411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                             RESULT 175
US-09-422-978-6039/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 176
US-09-371-772B-1411
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1.0%; Score 13.2; DB 1; Length 20; larity 83.3%; Pred. No. 1.9e+02; Conservative 0; Mismatches 3; Indels
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MEDIUW TYPE: EIDPOPY disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-DOS/MS-DOS
SOFTWARE: PATENTIN PATA:
APPLICATION NUMBER: US/O/982,759F
FILING DATE: 08-MAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9127319.3
FILING DATE: 23-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9221587.0
FILING DATE: 14-OCT-1992
ATTORNEY/AGENT INFORMATION:
NAME: BAKER, HOLLIE L.
REGISTRATION NUMBER: 31,321
REFERENCE/DOCKET NUMBER: 31,321
REFERENCE/DOCKET NUMBER: 102378.120
TELECOMMUNICATION INFORMATION:
TELEPRAK: 617-526-6000
APPLICANT: CZAPLEWSKI, Lloyd George
APPLICANT: GLIBERT, Richard
TITLE OF INVENTION: Stem Cell Inhibiting Proteins
NUMBER OF SEQUENCES: 178
CORRESPONDENCE ADDRESS:
ADDRESSEE: HALE and DORR LLP
STREET: 60 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature;
COATION: 1..20
CTHER INFORMATION: /product= "BB9513 oligomer"
0S-07-982-759F-134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Draper, Kenneth G.
APPLICANT: Draper, James
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: NF-KB
NUMBER OF SEQUENCES: 830
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-291-932A-311
; Sequence 311, Application US/08291932A
; Patent No. 5658780
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                820 GTCCTGATGCAGCTGAAG 837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 134:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 GIGCIGACGCAICIGAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA
                                                                                                                                                                 Boston
                                                                                                                                                                 STATE: MA
STATE: MA
7. 02109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       WEDIUM TYPE: Growing MEDIUM TYPE: Growing MEDIUM TYPE: BM PC Compatible COMPUTER: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APELICATION DATA: APPLICATION NUMBER: US/08/450,905B FILING DATE: 26-MAR-1995
FILING DATE: 08-MAR-1995
FILING DATE: 07/982,759
FILING DATE: 07/982,759
FILING DATE: 22-DEC-1991
PRIOR APPLICATION NUMBER: GB 9127319.3
FILING DATE: 21-DEC-1991
PRIOR APPLICATION NUMBER: GB 9221587.0
FILING DATE: 14-OCT-1992
ATTORNEY/AGENT INPORMATION:
NAME: NAME: DATE: 100071777
                                                                                                                                                                                                                                               Stem Cell Inhibiting Proteins 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature;
COCATION: 1..20
CHER INFORMATION: /product= "BB9513 oligomer"
198-08-95B-134
                                                                                               RESULT 179
US-08-450-905B-134/C
Sequence 114, Application US/08450905B
Fatent NO. 586501
GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: Stem Cell Inhibi
NUMBER OF ENDENCES: 178
CORRESPONDENCE ADDRESS:
ADDRESSEE: HALE and DORR
STREET: 60 State Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 180
US-07-982-759F-134/c
'S-equence 134, Application US/07982759F
') Patent No. 6057123
') GENERAL INFORMATION:
'APPLICANT: GRAIG, Stewart
'APPLICANT: GEORGE, Michael
') APPLICANT: EDWARDS, Richard Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: BAKER, HOLLE L.
REGISTRATION NUMBER: 31,321
REFERENCE/DOCKET NUMBER: 102,321
TELECOMMUNICATION INFORMATION:
TELEFAX: 617-526-610
INFORMATION FOR SEQ ID NO: 134:
SEQUENCE GHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDENESS: single
    1053 CAGCCCTGGCCTTCCCAT 1070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            820 GTCCTGATGCAGCTGAAG 837
                             1 CAGGCCUGACCUUCGCAU 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                          STATE: DC.
STATE: MA
02109
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Gaps

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Gaps

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Sequence 23, Application US/08250740
Patent No. 5686240
GENERAL INFORMATION:
APPLICANT: Schuchman, Edward H.
APPLICANT: Desnick, Robert J.
TITLE OF INVENTION: Acid Sphingomyelinase Gene and Diagnosis
TITLE OF INVENTION: Of Niemann-Pick Disease
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                    Score 13; DB 1; Length 17;
Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 y Match 1.0%; Score 13; DB 1; Length 17;
Local Similarity 100.0%; Pred. No. 1.4e+02;
hes 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARES PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/250,740
FILING DATE: 27-MAx-1994
CLASSIFICATION: 435
ATTONREY/AGENT INFORMATION:
NAME: CONTEXT, LAURE A.
REGISTRATION NUMBER: 30742
REFERENCE/DOCKET NUMBER: 6923-038
TELEPRANICATION INFORMATION:
TELEPRANICE: (212) 790-9090
TELEPRANICE: (212) 790-9090
                                                                                                                                                                                                                                                         1.0%; Sco...
100.0%; Pred. No. ...
... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                   TELEFAX: (619) 455-5110
INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS: LENGTH: 17 base pairs TYPE: nucleic acid STRANDEDNESS: single
     (619) 455-5100
                                                                                                                                                                                                                                                                                                                                              589 CTGCCCCCACCA 601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            631 CTCCAGGAGCTCT 643
                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                     CTGCCCCCCACCA 14
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: linear; MOLECULE TYPE: DNA US-08-250-740-23
                                                                                                                                                                                                             1..17
                                                                                                                                                                                           NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U
                                                                                                                                                                                                             ; LOCATION:
US-08-152-313-111
                                                                                                                                                                                                                                                                                                                                                                                                                                                              JS-08-250-740-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                        FEATURE
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Matches
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US-08-152-313-111
; Sequence 111, Application US/08152313
; Patent No. 5561041
; GENERAL INFORMATION:
; APPLICANT: Sidransky, David
; TITLE OF INVENTION: ANGLEIC ACID MUTATION DETECTION BY
TITLE OF INVENTION: ANALYSIS OF SPUTUM
; NUMBER OF SEQUENCES: 128
; CORRESPONDENCE ADDRESS:
; STREET: 1880 Century Park East, Suite 500
; CITY: Los Angeles
; STREET: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 13; DB 1; Length 15;
Pred. No. 1.1e+02;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-DOS/MS-DOS
SOFTWARE: PATENTIN DATA:
APPLICATION NUMBER: US/08/152,313
FILING DATE: 12-NOV-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Wetherell, Ur., Ph.D., John R.,
REGISTRATION NUMBER: 31,679
REFERENCE/DOCKET NUMBER: PD-2912
TELECOMMUNICATION INFORMATION:
           NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/157
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPA: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 311:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1066 CCCATCAGGCAGG 1078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 CCCAUCAGGCAGG 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 92.33
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
US-08-291-932A-311
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JULIOL STATES THE STATE OF THE SEQUENCE OF APPLICATION US/08985162

| PATENT NO. 6057156
| GENERAL INFORMATION:
| APPLICANT: Akhtar, Saghir APPLICANT: Akhtar, Saghir APPLICANT: Akhtar, Saghir TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT TITLE OF INVENTION: TO LEVELS OR CONDITIONS RELATED TITLE OF INVENTION: TO LEVELS OF EPIDERMAL GROWTH TITLE OF INVENTION: 1877
| CORRESPONDENCE ADDRESS: 1877
| CONTRESS OF EPIDER SEASON OF TO SEASON OF TAKEN OF THE SEASON OF TH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 17;
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                                                               COMPUTER REALIZED FORCE:

MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/695,472B
FILING DATE: 19910503
CLASSIFICATION: 435
ATTONBY/AGENT INPORMATION:
NAME: MISTOCK, S. Lealie
REGISTRATION NUMBER: 18,972
REGISTRATION NUMBER: 18,972
REGISTRATION NUMBER: 6923-014
TELEPHONE: (212) 790-9090
TELEPHONE: (212) 790-9090
TELEPRAN: (212) 790-9090
TELEFRAN: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.0%; Score 13; DB 1; Le
100.0%; Pred. No. 1.4e+02;
tive 0; Mismatches 0;
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Best Local Similarity 100.0
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: NUCLEIC ACID
STRANDEDNESS: single
TOPOLOGY: unknown
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TOPOLOGY: unknow;
; MOLECULE TYPE: DNA
US-07-695-472B-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-985-162-4/c
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0
                                                                                                   Sequence 111, Application US/08579223
; Patent No. 572609
; GENERAL INFORMATION:
; TITLE OF INVENTION: NUCLEIC ACID MUTATION DETECTION BY TITLE OF INVENTION: ANALYSIS OF SPUTUM
; VORRESPONDENCE ADDRESS:
; STREET: 1880 Century Park East, Suite 500
; CITY: Los Angeles
; TATE: California
; TATE: USA Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-07-695-472B-29

Sequence 29, Application US/07695472B

Sequence 29, Application US/07695472B

PRICENT NO. 5773278

GENERAL INFORMATION:

APPLICANT: Besnick, Robert J.

TITLE OF INVENTION: The Acid Sphingomyelinase Gene and TITLE OF INVENTION: Diagnosis of Niemann-Pick Disease

NUMBER OF SEQUENCES: 36

CORRESPONDENCE ADDRESS:

ADDRESSEE: Pennie & Edmonds

STREET: 1155 Avenue of the Americas

CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.0%; Score 13; DB 1; Length 17;
100.0%; Pred. No. 1.4e+02;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CONDITY: USA
CONDITY: USA
CONDITY: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Elb FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/579,223
FILING DATE: 28-DBC-1995
CLIASSIFFACATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/152,313
FILING DATE: 12-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: Wetherell, Jr., Ph.D., John R., REGISTRATION NUMBER: 31,678
REGISTRATION FOR SEQ ID NO: 111:
SEQUENCE CHARACTERISTICS:
MANDER: WALCHERISTICS:
MANDER: WALCHERISTICS:
MANDER: MANDER: WALCHERISTICS:
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MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         589 CIGCCCCCCACCA 601
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 13; Conserv
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US-08-579-223-111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY:
                                                                                          US-08-579-223-111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE:
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Gaps

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| Sequence 13, Application US/08469802B
| Patent No. 5741645
| Patent No. 5741645
| GENBERAL INFORMATION:
| APPLICANT: Orr, Harry T.
| APPLICANT: Chung, Ming-vi
| APPLICANT: Chung, Ming-vi
| APPLICANT: Cagabbi, Huda Y.
| TITLE OF INVENTION: Gene Sequence for Spinocerebellar Ataxia
| Patent No. 5741645
| TITLE OF INVENTION: Type 1 and Method for Diagnosis
| NUMBER OF SEQUENCES: 47
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Mueting, Raasch, Gebhardt & Schwappach, P.A.
| STREET: 119 No. 5741645th Fourth Street, Suite 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                       Sequence 111, Application PC/TUS9412947A
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES: 128
CORRESPONDENCE ADDRESS:
ADDRESSEE: Spensley Horn Jubas & Lubitz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 90067
COMPUTER READABLE FORM:
MUDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/12947A
FILING DATE: 10-NOV-1994
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
TELEFROME: (619) 455-5100
FELEFRAX: (619) 455-5100
INFORMATION FOR SEQ ID NO: 111:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TUDENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.0%; Score 13; DB 1; Le
100.0%; Pred. No. 1.4e+02;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                             ADDRESSEE: Spensley Horn Jubas & Lubitz
STREET: 1880 Century Park East, Suite 500
CITY: Los Angeles
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: DNA (genomic)
         631 CICCAGGAGCICT 643
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                                                  CICCAGGAGCICI 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.0
Matches 13; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
PCT-US94-12947A-111
                                                                                                                                          PCT-US94-12947A-111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 189
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100.0%; Pred. No. 1.4e+02;
tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 29, Application US/09106375
Patent No. 6541218
GENERAL INFORMATION:
APPLICANT: Schuchman, Edward H.
APPLICANT: Desnick, Robert J.
TITLE OF INVENTION: The Acid Sphingomyelinase Gene and TITLE OF INVENTION: Diagnosis of Niemann-Pick Disease NUMBER OF SEQUENCES: 36
CORRESPONDENCES. 36
STREEF: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: New York
STAIR: New York
COUTRY: U.S.A.
ZIP: 1036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Patentin PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRYT APPLICATION NUMBER: US/09/106,375
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ALIONAL FASENT ANTONIAL STATEMENT OF THE REGISTRATION NUMBER: 18,872
REPERENCE/DOCKET NUMBER: 6923-014
TELEPHONE: (212) 790-9090
TELEPHONE: (212) 790-9090
TELEX: 66141 PENNIE
TELEX: 66141 PENNIE
TELEX: 66141 PENNIE
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/695,472
FILING DATE: 03-MAX-1991
ATTORNEY/AGENT INFORMATION:
                  REFERENCE/DOCKET NUMBER: 230/107
TELECOMOUNICATION INFORMATION:
TELEFAX: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: (7-3510
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                200 CGGACGCCGACGA 212
                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.
Matches 13; Conservative
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                                                                                                                                                                     LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                             TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-106-375-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-106-375-29
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Length 18;
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1.0%; Score 13; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: BOBCOLL
STATE: BOBCOLL
STATE: MA
ZIP: 02109
COMPUTER READABLE FORM:
MEDING TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/450,905B
FRIGHT APPLICATION NUMBER: 07/982,759
FRIGH APPLICATION NUMBER: 07/982,759
FRILNG DATE: 30-DEC-1991
FRILNG DATE: 33-DEC-1991
FRILNG DATE: 33-DEC-1991
FRILNG DATE: 14-OCT-1992
ATTORNEY/AGENT INFORMATION:
NAME: BAKER, HOLLIE L.
REGESSATION NUMBER: 102.378.120DV-2
FILING DATE: 14-OCT-1992
ATTORNEY/AGENT INFORMATION:
NAME: BAKER, HOLLIE L.
RESTREMENCE/DOCKET NUMBER: 102.378.120DV-2
TELEPHONE: 617-526-610
TELEPHONE: 617-526-610
TELEPRAX: 617-526-610
SPONTENTY FRARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 191
US-08-450-905B-135/C
Sequence 135, Application US/08450905B
Fatent No. 5856301
GENERAL INFORMATION:
APPLICANT:
TITLE OF INTERFORMATION: Stem Cell Inhibiting Proteins
NUMBER OF SEQUENCES: 178
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                   1.0%; Score 13; DB 1; Le
100.0%; Pred. No. 1.7e+02;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; NAME/KEY: misc feature; LOCATION: 1..18 i OTHER INFORMATION: /product= "BB9516 oligomer" US-08-450-905B-135
                              information for SEQ ID NO: 3 SEQUENCE CHARACTERISTICS: LENGTH: 18 base pairs: TYPE: nucleic acid STRANDEDNESS: single: TOPOLOGY: linear MOLECTLE TYPE: DNA US-08-267-803B-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: HALE and DORR
STREET: 60 State Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                         592 CCCCCCACCAGCC 604
                                                                                                                                                                                                                                                                               13; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                     Best Local Similarity
Matches 13; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
                TELEFAX:
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Patent No. 5834183

GENERAL INFORMATION:

APPLICANT: Orr, Harry T.

APPLICANT: Chung, Ming-yi

APPLICANT: Chung, Ming-yi

TILLE OF INVENTION: Gene Sequence for Spinocerebellar Ataxia

TILLE OF INVENTION: Type 1 and Method for Diagnosis

NUMBER OF SEQUENCES:

CORRESPONDENCE ADDRESS:

ADDRESSEE: Mueting, Raasch, Gebhardt & Schwappach, P.A.

STREET: P.O. Box 581415

CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
1.0%; Score 13; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 13; Conservative 0; Mismatches 0; Indels
                                                                             COMPUTER FRADABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,802B
FILING DATE: 06-JUN-1995
CLASSTPICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: Mueting, Ann M.
REGISTRATION NUMBER: 110.00030101
TELECOMMUNICATION INFORMATION:
TELEBRANCE, DOS-1225
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: MN
COUNTRY: USA
ZIP: 55458-1415
COMPUTER READABLE FORM:
MEDLIOW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  110.00030120
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FILING DATE: 28-JUN-1994
CLASSIPICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MCCOTMACK, Myra H.
REGISTRATION NUMBER: 36,602
REFERENCE/DOCKET NUMBER: 110.00030120
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-305-1217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       592 CCCCCCACCAGCC 604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 190
US-08-267-803B-31/c
                                                            55401
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                                          COUNTRY:
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Gaps

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1140 TGCCTTTTTTTCT 1152
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             SEQUENCE CHARACTERISTICS.
LENGTH: 18 base pairs
TYPE: nucleic acid
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 100.
Matches 13; Conservative
                                                                                        single
                                                                                                                                                                                                                                                                                                                                                                                                         14 ATGTTGCTGACTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: primer_bind LOCATION: 1..18
                                                                                                        TOPOLOGY: linear MOLECULE TYPE: DNA
                                                              TYPE: nucleic
STRANDEDNESS:
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                                                                                                                                                                                                                                  US-07-982-759F-135
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                                                                                                                                                Sequence 29, Application US/09205860
Extent No. 5981732
GENERAL INFORMATION
FILE REPERBRORS:
CURRENT APPLICANT: US/09/205,860
CURRENT FILING DATE: 1998-12-04
NUMBER OF SEQ ID NOS: 87
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .;
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1.0%; Score 13; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: BOSTON
STATE: MA
STATE: MA
ZITY: BOSTON
STATE: MA
ZITY: 0.2109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/O7/982,759F
FILING DATE: 0.8-MAR-1993
RRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9127319.3
FILING DATE: 12-DEC-1991
PRIOR APPLICATION NUMBER: GB 9221587.0
FILING DATE: 14-OCT-1992
ATTORNEY/AGENT INFORMATION:
NAME: BAKER, HOLLIE L.
REGISTRATION NUMBER: 31,321
REPERBENCE/DOCKET NUMBER: 102378.120
TELEBCOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: CRAIG, Stewart
APPLICANT: CRAIG, Michael
APPLICANT: EDWARDS, Richard Mark
APPLICANT: CZAPLEWSKI, Lloyd George
APPLICANT: GILBERT, Richard
TITLE OF INVENTION: Stem Cell Inhibiting Proteins
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Antisense Oligonucleotide US-09-205-860-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 193
US-07-982-759F-135/c
; Sequence 135, Application US/07982759F
; Patent No. 6057123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: HALE and DORR LLP
STREET: 60 State Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Artificial Sequence
           758
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                                   14 ATGTTGCTGACTT 2
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        746 ATGITGCTGACTT
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TELEFAX: 6
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Sequence 8784, Application US/09422978

| Sequence 8784, Application US/09422978
| Patent No. 6537751
| GENERAL INNORMATION:
| APPLICANT: Chumakov, Daniel APPLICANT: Blumenfeld, Marta APPLICANT: Chumakov, Ilya ITLE REPERENCE: GENSET.020CP1
| TITLE NORMATION: Biallelic markers for use in constructing a high density...
| FILE REPERENCE: GENSET.020CP1
| CURRENT APPLICATION NUMBER: US/09/422,978
| GURRENT FILING DATE: 1999-04-21
| EARLIER PILING DATE: 1999-04-21
| EARLIER PILING DATE: 1999-1-23
| EARLIER PILING DATE: 1998-11-23
| EARLIER PILING DATE: 1998-11-23
| EARLIER PILING DATE: 1998-11-34
| WUMBER OF SEQ ID NOS: 11796
| SEQ ID NO 8784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: 1..18 -
OTHER INFORMATION: downstream amplification primer 99-18221 for SEQ 919, in comple
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GENERAL INFORMATION:
APPLICANT: Matteucci, Mark D.
APPLICANT: Matteucci, Mark D.
APPLICANT: Matteucci, Mark D.
TITLE OF INVENTION: CROSSLINKING AGENTS WHICH BIND TO THE MAJOR GROOVE OF TITLE OF INVENTION: DUPLEX DNA
TITLE OF INVENTION: DIPLEX DNA
TITLE OF INVENTION: DIPLEX DNA
NUMBER OF SEQUENCES: 158
CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster
STREET: 545 Middlefield Road, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                          Length 18;
                                                                                                                                              Query Match
1.0%; Score 13; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 13; Conservative 0; Mismatches 0; Indels
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NAME/KEY: misc feature
LOCATION: 1..18
OTHER INFORMATION: /product= "BB9516 oligomer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.0%; Score 13; DB 1; Le
100.0%; Pred. No. 1.7e+02;
iive 0; Mismatches 0;
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PCT-US91-03680-5
                          COUNTRY:
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GENERAL INFORMATION:
APPLICANT: Matteucci, Mark D.
APPLICANT: Matteucci, Mark D.
APPLICANT: Krawczyk, Steven
TITLE OF INVENTION: SEQUENCE-SPECIFIC NONPHOTOACTIVATED
TITLE OF INVENTION: DUPLEX DNA
TITLE OF INVENTION: DUPLEX DNA
NUMBER OF SEQUENCES: 158
CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster
STREET: 545 Midlefield Road, Suite 200
CITY: Menlo Park
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
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1.0%; Score 13; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 415-327-250
TELEFAX: 415-327-2951
TELEFAX: 415-327-2951
TELEFAX: 415-327-2951
TELEFAX: 415-327-2951
TELEFAX: 415-327-2951
TELEFAX: 416-327-2951
TENGTH: 18 base pairs
TYPE: NUCLEIC ACID
STRANDEDNES: single
TOPOLOGY: linear
                                                                                              COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US91/03680
FILING DATE: 19910524
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHBR
/note= "5-methylcytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: /mod base= OTHER OTHER INFORMATION: /mote= "5-methylcytosine' FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: 18
OTHER INFORMATION: /mod_base= OTHER
PCT-US91-03680-4
                                                                                                                                                                                                                                                                                                                                                                                                          4610-0011,40
                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: MATEANIGE, Kate H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 4610
TELECOMUNICATION INFORMATION:
TELEPHONE: 415-327-2550
TELEFAK: 415-327-2551
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EATURE:
NAME/KEY: modified_base
LOCATION: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1144 TITITITITIT 1156
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  Menlo Park
California
Y: USA
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OTHER INFORMATION:
OTHER INFORMATION:
                                                                              94025
CITY: Me
STATE: C
COUNTRY:
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COMPUTER READLE FORM:

WEINTY TYPE: TOPPY disk

COMPUTER: LIBR PC COMPATIBLE

ATTOMNY/AGENT INFORMATION:

FILLING DATE: WAREHING ATE H. 1.

REGISTRATION NUMBER: 23,359

TIELERHORM: 411-22-231

TIELERHORM: 411-22-231
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: Bavco, Pam
APPLICANT: APPLICANT: Stinchcomb, Dan
APPLICANT: Broobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
FILE REFERENCE: MBHB00, 97-1 (237/198)
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SECTION 6.925
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APPLICANT: Stinchoumb, Dan T.
APPLICANT: Draper, Kenneth
APPLICANT: Moswiggen, James
APPLICANT: Maryin, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 90071
COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/373,124A
                                                                                                                                                                                                                                                      Sequence 5925, Application US/09371772B Patent No. 6566127
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-373-124A-420
; Sequence 420, Application US/08373124A
; Patent No. 5646042
; GENERAL INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1311 GTAGCCAGGTGCTTTT 1326
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1 GGAGCCAGCUGCUUUU 16
                                         1 GTCTGCUGAGCGGGCU 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                             US-09-371-772B-5925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-371-772B-5925
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APPLICANT: Lochrie, Michael
APPLICANT: Joshic, Nebojsa
APPLICANT: Janjuc, Nebojsa
APPLICANT: Gold, Larry
TITLE OF INVENTION: Nucleic Acid Ligands Which Bind to Hepatocyte Growth
TITLE OF INVENTION: Factor/Scatter Factor (HGF/SF) or its Receptor C-Met
CURRENT APPLICATION NUMBER: US/09/364,539B
CURRENT FILING DATE: 1999-07-29
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LEMBER OF SEQ ID NOS: 192
SOFTWARE: Patentin Ver. 2.0
ERQ ID NO 10
LEMBER OF SEC ID NOS: 105
LEM
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NAME/KEY: modified base
LOCATION: (1)..(16)
COTHER INFORMATION: Purines and pyrimidines are 2'OMe, purines and
OTHER INFORMATION: pyrimidines at positions 1-4 are DNA; purines and
OTHER INFORMATION: pyrimidines at positions 5-16 are RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%; Score 13; DB 1; Length 18;
100.0%; Pred. No. 1.7e+02;
tive 0; Mismatches 0; Indels
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/04094
FILING DATE: PCT/US95/04094
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/224,010
FILING DATE: 06-APR-1994
ATFORNEY/AGENT INPORMATION:
NAME: Crane-Feury, Sharon E
REGISTRATION NUMBER: 36,113
REGISTRATION NUMBER: 36,113
REGISTRATION NUMBER: 36,113
REGISTRATION NUMBER: 36,113
REGISTRATION STATE OF STATE O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 198
US-09-364-539-10
; Sequence 10, Application US/09364539B
; Patent No. 6344321
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               297 GTCTGCTGTGGGGCT 312
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Artificial Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                379 CTTCCTCCAGAGG 391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
MOLECULE TYPE: DNA (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best_Local Similarity
Matches 13; Conserva
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US-09-364-539-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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Gaps
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Pred. No. 1.6e+02;
6; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.9%; Score 12.8; DB 1;
Best Local Similarity 75.0%; Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-78-306-655

Sequence 655, Application US/08758306

Patent No. 5807743

Patent No. 5807743

Patent No. 5807743

APPLICANT: Stinchcomb, Dan T.

APPLICANT: Stinchcomb, Dan T.

APPLICANT: McSwiggen, James A.

TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: TREATMENT OF DISBASES

TITLE OF INVENTION: ASSOCIATED WITH

TITLE OF INVENTION: ASSOCIATED WITH

TITLE OF INVENTION: GAMMA-CHAIN EXPRESSION

NUMBER OF SEQUENCES: 1379

CORRESPONDENCE ADDRESS:

ADDRESSE: Lyon & Lyon

STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: 1BM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSeq Version 1.5
CURRENT APPLICATION DATA:
APPLICATION DATE: DECEMBER: US/1996
FILING DATE: December 3, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 212/132
TELECOMOMULATORION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1095 TGAACGTAATTATGTA 1110
                                           TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2031
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.9%;
Best Local Similarity 50.0%;
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: 67.3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                    ; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-373-124A-2031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ð
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 12.8; DB 1; Length 17; 62.5%; Pred. No. 1.6e+02; ive 4; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INCEMENTION:

APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Draper, Kenneth
APPLICANT: McWiggen, James
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: TREATMENT OF RESTRUCIS AND
TITLE OF INVENTION: TREATMENT OF RESTRUCIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: G33 West Fifth Street
CONNENT: Los Angeles
COTY: Los Angeles
CONNENT: U.S.A.
ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: SLOCAGE
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: Mord Perfect 5.1
CURRATING SYSTEM: IBM P.C. DOS 5.0
SOFTWART APPLICATION MIMBED: 10.000
## PILING DATE: February 7, 1994
## APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
## FILING DATE: Agust 26, 1992
## ATTORNEY AGENT INFORMATION:
## ATTORNEY AGENT INFORMATION:
## ATTORNEY AGENT INFORMATION:
## REGISTRATION NUMBER: 32,327
## REGISTRATION NUMBER: 299/035
## REGISTRATION NUMBER: 299/035
## REGISTRATION INFORMATION:
## TELEFHONE: (213) 489-1600
## TELEFAX: (213) 955-0440
## TELEFAX: (213) 955-044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/373,124A FILING DATE: January 13, 1995
PRIOR APPLICATION DATA: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2031, Application US/08373124A Patent No. 5646042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
FILING DATE: AUGUST 26, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     795 CCTGGCTCGCTCCTG 810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 ccudgcucccuaccug 17
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Best Local Similarity 62.5°
Matches 10, Conservative
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Length 17;
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              APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.9%; Score 12.8; DB 1;
62.5%; Pred. No. 1.6e+02;
tive 4; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,628
FILING DATE: 05-MAY.1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/373,124
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/245,466
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/943,7111NG DATE: Pebruary 7, 1994
APPLICATION NUMBER: 07/943,7111NG DATE: December 7, 1994
APPLICATION NUMBER: 07/943,7111NG DATE: December 7, 1992
APPLICATION NUMBER: 07/943,7132
FILING DATE: August 26, 1992
ATTONNEY, AGENT INFORMATION:
NAME: WARDLEY, RICHARD
REGISTRATION NUMBER: 07/936,422
ATTONNEY, AGENT INFORMATION:
NAME: WARDLEY, RICHARD
                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2031, Application US/08435628
; Patent No. 5817796
; GENERAL INFORMATION:
    APPLICANT: Stinchcomb, Dan T.
    APPLICANT: McSwiggen, James
    APPLICANT: Jaryls, Thale
    TITLE OF INVENTION: METHODS AND COMPO
; TITLE OF INVENTION: TREATMENT OF REST
; TITLE OF INVENTION: CANCER USING RIBO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 209/035
TELECOMMULICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELERAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 795 CCTGGCTCGCTCCTG 810
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                              STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 10; Conserv
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US-08-435-628-2031
                                                                                                                                                                                                                                                                                              90071
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Gaps
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0.9%; Score 12.8; DB 1; Length 17;
Best Local Similarity 81.2%; Pred. No. 1.6e+02;
Matches 13; Conservative 1; Mismatches 2; Indels
  2; Indels
                                                                                                                                                                                                                                         GENERAL NO. 30.713.

GENERAL NO. 30.713.

APPLICANT: Stinchcomb, Dan T.

APPLICANT: Stinchcomb, Dan T.

APPLICANT: Stinchcomb, Dan T.

APPLICANT: Stinchcomb, Dan T.

TITLE OF INVENTION: METHOD NITH

TITLE OF INVENTION: TREATMENT OF DISEASES

TITLE OF INVENTION: INTERLEUKIN-2 RECEPTOR

TITLE OF INVENTION: GAMMA-CHAIN EXPRESSION

NUMBER OF SEQUENCES: 1379

CORRESPONDENCE ADDRESS:

ADDRESSE: ADDRESSE:

STREET: 633 West Fifth Street
  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FrattSeq Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/758,306
FILING DATE: December 3, 1996
CLASSIFICATION: 514
PRICE APPLICATION: 104
APPLICATION NUMBER: 212/13
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 212/132
REFERENCE/DOCKET NUMBER: 212/132
TELECOMMUTICATION INFORMATION:
TELEPHONE: (213) 955-0440
TELEPAX: (213) 955-0440
TELEPAX: 67-3510
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-435-628-420; Sequence 420, Application US/08435628; Patent No. 5817796; GENERAL INFORMATION: APPLICANT: Stinchcomb, Dan T.; APPLICANT: Draper, Kenneth
                                                                                                                                                                              US-08-758-306-721
; Sequence 721, Application US/08758306
; Patent No. 5807743
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                                               227 CTCAGCCTCAGGCATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
  12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Los Angeles
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-758-306-721
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: L
STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 204
  Matches
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                                                                                                                                                            MEDIUM TYPE: storage
COMBUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620A
FILNG DATE: Adgust 17, 1994
CLASSIFICATION DATA: Hockeribed below:
PRIOR APPLICATION DATA: Gescribed below:
PRIOR APPLICATION DATA: 10, 1993
PRIOR APPLICATION DATA: 10, 1993
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 03/207
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 208/149
TELECOMMUNICATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 955-0440
TELEPHONE: (213) 955-0440
TELEPHONE: G7-3510
INFORMATION FOR SEQ ID NO: 1727:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: mucleic acid
STRANDENDESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-29-620A-1937/C
US-08-29-620A-1937/C
IS-08-29-620A-1937/A-2PDIcation US/08292620A
| Sequence 1937, Application US/08292620A
| Patent No. 5837542
| APPLICANT: Susan Grimm
| APPLICANT: Dan T. StinchComb
| APPLICANT: Dan T. StinchComb
| APPLICANT: Sean Sullivan
| APPLICANT: Remeth G. Draper
| TITLE OF INVENTION: RELATED TO LEVELS OF
| TITLE OF INVENTION: RELATED TO LEVELS OF
| TITLE OF INVENTION: INTACELLULAR ADHESION
| TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
| NUMBER OF SEQUENCES: 2390
| CORRESPONDENCE ADDRESSE:
| ADDRESSEE: Lyon & Ive-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700 CITY: Los Angeles STATE: California COUNTRY: U.S.A. ZIP: 9071-2066 COMPUTER READABLE FORM: MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                 COUNTRY: U.S.A.
ZIF: 90071-2066
COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 334 CCTGGTGATAGTCACA 349
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Suite 4700
                      CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
TOPOLOGY: 1:
US-08-292-620A-1727
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50.0%; Pred. No. 1.6e+02;
tive 6; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-22-620A-1727/c
Sequence 1727, Application US/08292620A
Patent No.: 5837545.
GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwiggen
APPLICANT: Sean Stlinchcomb
APPLICANT: Sean Sullivan
APPLICANT: Seneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: NOLECULE-1 (I-CAM-1)
                                                                                                                                                       COUNTRY: U.S.A.

ZIP: 90071
ZIP: 90071
ZIP: 90071
ZIP: 90071
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: Ward Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,628
FILING DATE: 05-MAY-1995
CLASSIFICATION NUMBER: 08/373,124
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/345,466
FILING DATE: February 7, 1994
APPLICATION NUMBER: 08/245,466
FILING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 08/32,943
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/984,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
APPLICATION NUMBER: 07/936,422
APPLICATION NUMBER: 209/035
FILING DATE: BECOMMETION:
CELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEST: 67-3510
TELEST: 67-3510
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633 West Fifth Street
       CORRESPONDENCE ADDRESS:
ADDRESSEB: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1095 TGAACGTAATTATGTA 1110
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 50.03
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                  CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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STREET: 63
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Patent No. 6057156
GENERAL INFORMATION:
APPLICANT: Achtar, Saghir
APPLICANT: Fall, Patricia
APPLICANT: McSwiggen, James
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT:
TITLE OF INVENTION: TO LEVELS OF ENDERMAL GROWTH
TITLE OF INVENTION: TO LEVELS OF ENDERMAL GROWTH
TITLE OF SEQUENCES: 1877
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.9%; Score 12.8; DB 1; Best Local Similarity 87.5%; Pred. No. 1.6e+02; Matches 14; Conservative 0; Mismatches 2;
                                                                                    NAME: Murashige, Kate H
REGISTRATION NUMBER: 29,959
REGISTRATION NUMBER: 35029-20001.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-887-1500
TELEPAX: 202-822-0168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIE: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: Storage
COMPUTER: Factage
COMPUTER: Factage
COMPUTER: BM Compatible
COMPUTER: Factage
COMPUTER: BM Compatible
COMPUTER: BM Compatible
COMPUTER: Factage
COMPUTER: BM Compatible
COMPUTER: Factage
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NAME/KEY: Other

LOCATION: 1...17

OTHER INFORMATION: HIP sequence
US-08-765-783A-79
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STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        870 CCCCACAGCCAAGTTC 885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 cccchahecchheric 17
                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-985-162-452
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 209
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MEDIUM TYPE: storage
COMPUTER: IBM Compatible
SOFTWARE: Word Perfect 5.1
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
FILING DATE: AQUAGE: 17, 1994
CLASSIFICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION NUMBER: 08/008/895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGRAT INCORMATION:
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELEFARICATION INFORMATION:
TELEFAN: 67-3510
TELEFAX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: MORRISON & FORESTER
STREET: 2000 Pennsylvania Avenue, NW, suite 5500
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-765-783A-79

'Sequence 79, Application US/08765783A

'Sequence 79, Application US/08765783A

'Sequence 79, Application US/08765783A

'Settent No. 294524

'APPLICANT: Matsushima, Kouji

APPLICANT: Matsusda, Yoshiki

APPLICANT: Sato, Koh

APPLICANT: Tsuchiya, Masayuki

APPLICANT: Tsuchiya, Masayuki

APPLICANT: Tsuchiya, Masayuki

APPLICANT: Tsuchiya, Masayuki

TITLE OF INVENTION: Interleukin-8

NUMBER OF SEQUENCES: 105

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/765,783A
FILING DATE: 07-MAR-1997
CLASSIFICATION DATA:
APPLICATION DATA:
FILING DATE:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Washington
STATE: DC
COUNTRY: USA
ZIP: 20006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            334 CCTGGTGATAGTCACA 349
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US-08-292-620A-1937
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Best Local Similarity 87.5%; Pred. No. 1.6e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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APPLICANT: Susan Grimm
APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: Gens Sullivan
APPLICANT: Sean Sullivan
APPLICANT: Renneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: NURACELLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2330
CORRESPONDENCE: 2330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IEBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE:
CLASSIFICATION NUMBER: US/09/071,845
FILING DATE:
APPLICATION NUMBER: US/08/292,620
FILING DATE: AUGUST 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 08/008,895
FILING DATE: Jenember 7, 1992
APPLICATION NUMBER: 07/989,849
RTLING DATE: December 7, 1992
ATTORNEY AGENT INFORMATION:
NAME: WATCHLY RICHART ON MUMBER: 32,337
RECISTRATION WUMBER: 32,337
REFERENTE OF THE TABLE OF TABLE OF THE TABLE OF THE TABLE OF THE TABLE OF THE TABLE OF TABLE OF TABLE OF THE TABLE OF TA
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STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: SLORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-071-845-1937/c
; Sequence 1937, Application US/09071845
; Patent No. 6132967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 495-1600
TELEPRAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                 334 CCTGGTGATAGTCACA 349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         334 CCTGGTGATAGTCACA 349
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SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 87.5 Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-071-845-1937
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ALESULT 210

US-09-071-845-1727/C

Sequence 1727, Application US/09071845

Patent No. 6132967

GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RIBOZYME TO LEVELS OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: NOLECULE-1 (I-CAM-1)
CORRESPONDENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: LYON & INVENTION
STREFT: ADDRESSEE: LYON & INVENTION
ATREED OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Brorage
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION TOWNER: US/09/071,845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      THILMS DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: Date North Nor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                 574 CAGCAGGCCCTCCGTC 589
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
         LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                     ; TOPOLOGY:
US-08-985-162-452
LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
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Gaps
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0
0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 1.6e+02; Live 0; Mismatches 2; Indels
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0.9%; Score 12.8; DB 1; Length 17;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: Pavco, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                           COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
COMPUTER: ENCAQUE COMPATION
SOFTWARE: WORD FOR S.0
SOFTWARE: WORD PEFfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: OCTOBER 26, 1995
ATTORNEY AGENT INFORMATION:
NAME: WASDLEY, RICHARD S. 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 7924, Application US/08584040; Patent No. 6346398; GENERAL INFORMATION:
              TITLE OF INVENTION: INCADENTIONS RELIGIOUS RELIGIONS TO THILE OF INVENTION: OF VASCULAR ENTITLE OF INVENTION: OF VASCULAR ENTITLE OF INVENTION: OF VASCULAR ENTITLE OF INVENTION: OF OWNERS.

CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 613 West Fifth Street
STREET: 614te 4700
CITY: Los Angeles
STREET: California
COUNTRY: U.S.A.
ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.9%;
Best Local Similarity 87.5%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17 CAAAATCTGAGCAGAC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41 CAAAATCTTAGCATAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: 1
US-08-584-040-4374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-584-040-7924
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                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 Pennsylvania Avenue, NW, suite 5500
STRYE: DC
                                                                                Parent No. 6245894
GENERAL INFORMATION:
APPLICANT: Mateushima, Kouji
APPLICANT: Mateushima, Kouji
APPLICANT: Mateumoto, Yoshihiro
APPLICANT: Sato, Koh
APPLICANT: Tsuchiya, Masayuki
APPLICANT: Tsuchiya, Masayuki
APPLICANT: Tsuchiya, Masayuki
TITLE OF INVENTION: Reshaped Human Antibody to
TITLE OF INVENTION: Interleukin-8
NUMBER OF SEQUENCES: 105
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'SGQUENCE 4374, Application US/08584040
'SGQUENCE 4374, Application US/08584040
'PATENT NO. 6346398
'APPLICANT: Pavco, Pamela
'APPLICANT: McSwiggen, James
'APPLICANT: Stinnchomb, Dan T.
'APPLICANT: Bscobedo, Jaime
'ITILE OF INVENTION: METHOD AND REAGENT FOR THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 20006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM COMPALIALE
OPERATING SYSTEM: DOS
SOFTWARE: FASISED for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/416,557
FILING DATE: 12-October-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         35029-20001.10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFCATION
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/765,783
FILING DATE: 7-March-1997
ATTORNEY/AGENT INFORMATION:
NAME: Murashige, Kate H
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 35029-27
TELEPHONE: 20-887-1500
TELEPHONE: 20-887-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: 1...17
OTHER INFORMATION: HIP sequence
US-09-416-557-79
                                                              ; Sequence 79, Application US/09416557; Patent No. 6245894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 870 CCCCACAGCCAAGTTC 885
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO. SEQUENCE CHARACTERISTICS: LENGTH: 17 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                               US-09-416-557-79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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Gaps

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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Mello, Fatthcha Ann Owens
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Folkerts, Otto
APPLICANT: Folkerts, Otto
APPLICANT: Mello, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR
TITLE OF INVENTION: MODULATION OF GENE EXPRESSION
TITLE OF INVENTION: IN PLANTS
NUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
COUNTY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: UNJY 12, 1996
CLASSIFICATION DATA:
APPLICATION WUMBER: 60/001,135
FILING DATE: UNJY 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: SEPTEMBER: 01994
                       CLASSIERCATION BOTA:
CLASSIERCATION 800

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/001,135
FILING DATE: 30,194
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: WAIDLIGY RICHARD 32,327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Zwick, Michael G.
APPLICANT: Edington, Brent E.
APPLICANT: McSwiggen, James A.
APPLICANT: Merlo, Patricia Ann Owens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 216
US-08-679-645-886
; Sequence 886, Application US/08679645
; Patent No. 6350934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               891 GCTGCGGTACAGCGTG 906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GCUGCGGUUCAGCCUG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 68.8<sup>†</sup>
Matches 11, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: SID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TOPOLOGY:
US-08-679-645-75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   à
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
0.9%; Score 12.8; DB 1; Length 17;
Best Local Similarity 56.2%; Pred. No. 1.6e+02;
Matches 9; Conservative 5; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 75, Application US/08679645

Sequence 75, Application US/08679645

GENERAL INFORMATION:
APPLICANT: Edington, Brent E.
APPLICANT: Edington, James A.
APPLICANT: McSwiggen, James A.
APPLICANT: McSwiggen, James A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Scott A.
APPLICANT: Scott A.
APPLICANT: Scott A.
APPLICANT: ONDG, Donald J.
ITLE OF INVENTION: COMPOSITION AND METHODS FOR TITLE OF INVENTION: IN PLANTS
NUMBER OF SEQUENCES: 1263
ADDRESSED: LYON & LYON
STREFF. LYON & LYON
STREFF.
COMPUTER REALBABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFTCATION 514
PRIOR APPLICATION DATE:
APPLICATION NUMBER: 60/005,974
FILING DATE: OCTOBER 26, 1995
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 26,1995
ATTORNEY/AGENT INFORMATION:
TELEROWEL/DOCKET NUMBER: 218/064
TELEROWEL/DOCKET NUMBER: 218/064
TELEROWEL/DOCKET NUMBER: 218/064
TELEROWEL/DOCKET NUMBER: 218/064
TELEROWEL/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELEROWEL/DOCKET NUMBER: 32,327
RELEROWER (213) 955-0440
INFORMATION FOR SED ID NO: 7924:
TELERX: 67-3510
INFORMATION FOR SED ID NO: 7924:
TELERX: 67-3510
INFORMATION FOR SED ID NO: 7924:
TELERX: 67-3510
INFORMATION FOR SED ID NO: 7924:
TELERY: 67-3510
INFORMATION FOR SED ID NO: 7724:
TELERY: 67-3510
INFORMATION FOR SED ID NO: 7724:
TELERY: 67-3510
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STATE: California
COUNTRY: U.S.A.
ZIP: 20071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
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STREET: 633 West Fifth Street
STREET: Suite 4700
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1 CUUCGCAUACUGCUCA 16
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US-08-584-040-7924
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GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Rayoro, Pam
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stochedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions:
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: 1999-108
TITLE OF INVENTION NUMBER: US 60/005, 974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-36
                                           APPLICANT: McSwiggen, Jim
APPLICANT: Strindcomb, Dan
APPLICANT: Strindcomb, Dan
APPLICANT: Strindcomb, Dan
APPLICANT: Extended and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1995-08-10
PRIOR PLING DATE: 1995-10-26
PRIOR PLING DATE: 1995-10-26
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin Version 3.0
LENGTH: 17
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US-003-31-72B-4175
US-00-371-72B-4175
Sequence 4175 Application US/09371772B
Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.; APPLICANT: PavCo, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 3707, Application US/09371772B Patent No. 6566127
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Best Local Similarity 87.5
Matches 14; Conservative
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US-09-371-772B-2141
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Best Local Similarity
Matches 9; Conserv
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US-09-371-772B-3707
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APPLICANT: Burgin, Alex
APPLICANT: Burgin, Alex
APPLICANT: Beaudry, Andex
APPLICANT: Beaudry, Andex
APPLICANT: Seedler, David
APPLICANT: Sweedler, David
APPLICANT: Sweedler, David
APPLICANT: Zinnen, Shawn
ITLE OF INVENTION: Nucleotide triphosphate and their incorporation into oligonucleot
GURRENT APPLICATION NUMBER: US 69/2474,432B
CURRENT FILING DATE: 1999-12-19
PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR PELING DATE: 1999-11-05
PRIOR PELING DATE: 1999-11-05
PRIOR PELING DATE: 1999-11-05
PRIOR PELING DATE: 1999-11-04
PRIOR FILING DATE: 1998-11-04
PRIOR PELING DATE: 1998-11-04
PRIOR PELING DATE: 1999-11-04
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81.2%; Pred. No. 1.6e+02;
tive 1; Mismatches 2;
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US-09-371-772B-2141/c
; Sequence 2.141, Application US/09371772B
; Setent No. 6566127
; GENERAL INFORMATION:
; APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 388, Application US/09474432B Patent No. 6528640.
GENERAL INFORMATION.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                219/247
                       NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REEECOMMICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: (7-3510
INFORMATION FOR SEQ ID NO: 886:
SEQUENCE CHRACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1103 ATTATGTAGTTTTCTG 1118
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2 AUTUTGUAUTUTCUG 17
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SOFTWARE: Patentin version 3.0
SEQ ID NO 388
LENGTH: 17
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ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 81.2
Matches 13; Conservative
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CORGANISM: Homo sapiens
US-09-474-432B-388
                                                                                                                                                                                                                                                                                                                                                                                                                linear
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                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY:
US-08-679-645-886
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US-08-219-842-85/C
) Sequence 85, Application US/08219842
) Sequence 85-65533
) GENERAL INFORMATION:
) APPLICANT: Parker, W. D.
) APPLICANT: Herristadt, Corinna
) TITLE OF INVENTION: Diagnostic and Therapeutic Compositions
) TITLE OF INVENTION: for Alzheimer's Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                       COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/219,842
FILING DATE: 30-MAR-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
REFERENCE/COCKET NUMBER: 31.815
REFERENCE/COCKET NUMBER: 32.815
INFORMATION FOR SEQ ID NO: 52.
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TELENGTH: 18 base pairs
                            NUMBER OF SEQUENCES: 95
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 95
CORRESPONDENCE ADDRESS:
ADDRESSEB: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DAIA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 12.8; DB 1;
Pred. No. 1.8e+02;
              FITLE OF INVENTION: for Alzheimer's Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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FILING DATE: 30-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
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NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31, 31, 315
REFERENCE/DOCKET NUMBER: P-A
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.9%;
Best Local Similarity 87.5%;
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STRANDEDNESS: single
                                                                                                  ALU...
STREET: 43,0...
CITY: San Diego
TITY: California
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STATE: California
COUNTRY: USA
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                                                                                                                                                                                        COUNTRY: US
ZIP: 92122
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                APPLICANT: StinchComb, Dam
APPLICANT: StinchComb, Dam
APPLICANT: StinchComb, Dam
APPLICANT: StinchComb, Dam
APPLICANT: StinchComb, Dame
TITLE CP INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PILING DATE: 1996-01-08
SOFTWARE PAILS DATE: 1996-01-08
SOFTWARE PAILS DATE: 1996-01-08
SOFTWARE PAILS DATE: 1996-01-08
SOFTWARE PAILS DATE: 1997-04-108
SEC ID NO 4175
LENGTH: 17
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APPLICANT: Stinchcomb, Dan
APPLICANT: Bescobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
FILE REFERENCE: MEHOO, 376-01
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
SEQ ID NO 6941
LENGTH: 17
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Patent NO. 5565323
GENERAL INFORMATION:
APPLICANT: Parker, W. D.
APPLICANT: Herrnstadt, Corinna
TITLE OF INVENTION: Diagnostic and Therapeutic Compositions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%; Score 12.8; DB 1; Length 17; 81.2%; Pred. No. 1.6e+02; rive 1; Mismatches 2; Indels
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Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              727 CAGGGGCCTGGCTGC 742
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Best Local Similarity 81.29
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
ORGANISM: Homo sapiens
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Best Local Similarity
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US-09-371-772B-6941/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-371-772B-4175
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US-08-219-842-52
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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US-08-451-096-85/c

Sequence 85, Application US/08451096

Fatent No. 5760205

GENERAL INFORMATION:

APPLICANT: Parker, W. D.

APPLICANT: Parker, W. D.

TITLE OF INVENTION: for Alzheimer's Disease

TITLE OF SEQUENCES: 95

CORRESPONDENCE ADDRESS:

ADDRESSEE: Campbell and Flores

STREET: 4370 La Jolla Village Drive, Suite 700

CITY: San Diego

STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .
0
                                                                                                                                                   Sequence 52, Application US/08451096
Patent No. 5760205
GENERAL INFORMATION:
APPLICANT: Parker, W. D.
APPLICANT: Herrnstadt, Corinna
TITLE OF INVENTION: Diagnostic and Therapeutic Compositions
TITLE OF INVENTION: Diagnostic and Therapeutic Compositions
NUMBER OF SEQUENCES: 95
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 1.8e+02; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPOURE READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATUR SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                 E: Campbell and Flores
4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-AG 9504
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/219,842
FILING DATE: 30-MAR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/451,096
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (619) 535-9001
TELEPAX: (619) 535-8949
INPORMATION FOR SEQ ID NO: 52:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
             924 GATGGCAGATCTGGAG 939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           590 TGCCCCCCACCAGCCT 605
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                                                    18 GGTGGCTGATCTGGAG 3
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Best Local Similarity 87.55
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                       STREET: 4370 La Joi
CITY: San Diego
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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                                                                                                                RESULT 225
US-08-451-096-52
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US-08-451-096-52
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                                                                                                                                                                                                           Query Match 0.9%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 1.8e+02; Matches 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                             KESULT 224

US-08-363-240A-1187/c

Sequence 1187, Application US/08363240A

Patent No. 5705388

GENERAL INFORMATION:
APPLICANT:
APPLICANT: Couture, Larry
APPLICANT: McSwiggen, James
APPLICANT: Bisgaier, Charles
APPLICANT: Bisgaier, Charles
APPLICANT: Pape, Michael

TITLE OF INVENTION: PREVENTION INHIBITION OF
TITLE OF INVENTION: PROFESSION
TITLE OF INVENTION: OF VASCULAR DISEASES
CORRESPONDENCE ADDRESS:
ADDRESSES: Lyon F T
STREET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A.
ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: World Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US/08/363,240A
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/363,2
FILING DATE: December 23, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 312,327
REFRENCE/DOCKET NUMBER: 210/I
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                               590 TGCCCCCACCAGCCT 605
  (619) 535-9001
                   TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 67.3510
INPORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                 LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
1 TOPOLOGY: linear
US-08-219-842-85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.9
Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-363-240A-1187
TELEPHONE:
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Gaps
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 580//50/
GENERAL INFORMATION:
APPLICANT: 1TO, Kiyoshi
APPLICANT: YAMAKI, Toshifumi
APPLICANT: ARIL, Teruo
APPLICANT: TSURUOKA, Miyuki
APPLICANT: NAKAMURA, Takeshi
TITLE OF INVENTION: NOVEL NITRILE HYDRATASE
                                                                                                                                                                                                                                                                                                                           Query Match 0.9%; Score 12.8; DB 1; Best Local Similarity 87.5%; Pred. No. 1.8e+02; Matches 14; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACCOUNTY: Alexandria
STREET: P.O. BOX 1404
CITY: Alexandria
CITY: Alexandria
STRATE: Virginia
COMPUTER: EDAPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/800,751
                            TELEPHONE: (703) 836-6620
TELETAX: (703) 836-6620
INPOMATION FOR SEQ ID NO: 39:
SEQUENCE: ARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDENESS: single
TOPOLLGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "synthetic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         , NOLECULE TYPE: other nucleic acid
; DESCRIPTION: /desc = "synthetic DNA"
US-08-800-751-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION: 435
FILING DATE:
APPLICATION DATE:
APPLICATION DATE:
FILING DATE:
TING DATE: 14-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 35,030
TELEPRONGIO TONEONATION:
TELEPRONGIO TONEONATION:
TELEPRONE: (703) 836-6620
TELEPRANE: (703) 836-2021
INFORMATION FOR SEQ ID NO: 40: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 228
US-08-800-751-40
; Sequence 40, Application US/08800751
; Patent No. 5807730
          TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                            879 CAAGTTCCAGGAGCTG 894
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid
EDNESS: single
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STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: li:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 39, Application US/08800751
Sequence 39, Application US/08800751
Patent No. 5807730
GENERAL INFORMATION:
APPLICANT: TAO, Kiyoshi
APPLICANT: TO, Kiyoshi
APPLICANT: TANENCA, Toshifumi
APPLICANT: TSURUOKA, Miyuki
APPLICANT: NAKAMURA, Takeshi
ITITLE OF INVENTION: NOVEL NITRILE HYDRATASE
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
STREET: P.O. Box 1404
STREET: P.O. Box 1404
COUNTRY: United States
STREET: Virginia
COUNTRY: United States
INDIM TYPE: Ploppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: IBM PC Compatible
OPERATING SYSTEM: Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
FILING DATE:
FILING DATE:
FILING DATE:
COUNTRY: USA

ZIP: 97122

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARR: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/451,096
                                                                                                                                                                                                               FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
PRIOR APPLICATION: BATA:
APPLICATION NUMBER: US 08/219,842
FILING DATE: 30-MAR-1994
ATTORNEY-AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: P-AG 9504
TELEFONMICATION INFORMATION:
TELEFONMICATION INFORMATION:
TELEFONMICATION INFORMATION:
TELEFONMICATION INFORMATION:
TELEFONMICATION SEQ ID NO: 855.
ENGURNE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
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REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 028022-007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   590 TGCCCCCCACCAGCCT 605
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ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear
US-08-451-096-85
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US-08-800-751-39
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SELECTION MARKER GENE FREE RECOMBINANT
STRAINS: A METHOD FOR OBTAINING THEM AND THE USE OF THESE
STRAINS
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        THEIR USE IN THERAPEUTIC AND DIAGNOSTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.9%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 1.8e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: SELTEN, GERARDUS CORNELIUS MAIA
APPLICANT: SMINKELS, BART WILLEM
APPLICANT: VAN GORCOM, ROBERTUS
APPLICANT: VAN GORCOM, ROBERTUS
APPLICANT: FRANCISCUS MARIA
TITLE OF INVENTION: SELECTION MARKER GENE FREE RECOMI
TITLE OF INVENTION: STRAINS: A METHOD FOR OBTAINING
TITLE OF INVENTION: STRAINS
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSE: MORRISON & FOERSTER
STREET: 2000 Pennsylvania Ave. N.W., Suite 5500
CITY: Washington, D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 2006-1888
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/880,557
FILING DATE: 23-UUN-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
TITLE OF INVENTION: THEIR USE IN THERAPES
TITLE OF INVENTION: METHODS
NUMBER OF SEQUENCES: 39
CORRESPONDENCE ADDRESS:
ADDRESSEB: MORGAN & FINNEGAN, L.L.P.
STREET: 345 PARK AVENUE
CITY: NEW YORK
STATE: NEW YORK
                                                                                                                                                                                                                      COMPUTER READABLE FORM:
WEDIUM TYPE: TEOPPY DISK
COMPUTER: FLOPPY DISK
COMPUTER: ISM PC COMPATIBLE
COMPUTER: ISM PC COMPATIBLE
COMPUTER: ISM PC COMPATIBLE
COMPAGE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/411,098
FILING DATE: 27-MAR-1995
CLASSIFFCATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: CAROL M. GRUPPI
REGISTRATION NUMBER: 37,341
REGISTRATION NUMBER: 37,341
REFERENCE/DOCKET NUMBER: 2026-4188
TELECHOME: (212) 758-4800
TELEFAX: (212) 758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2026-4188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-880-557-18

; Sequence 18, Application US/08880557

; Patent No. 5876988

; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELLEFAX: (212) 751-684
TELLEX: 421792
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: SINGLE TOPOLOGY: UNKNOWN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: NUCLEOTIDE
                                                                                                                                                                                                            USA
                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-411-098-35
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                                                                                                                                                                                         Sequence 971, Application US/08758306
Fatent No. 5807743
GENERAL INFORMATION:
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: TREATMENT OF DISEASES
TITLE OF INVENTION: TREATMENT OF DISEASES
TITLE OF INVENTION: INTERLEUKIN-2 RECEPTOR TITLE OF INVENTION: GAMMA-CHAIN EXPRESSION NUMBER OF SEQUENCES: 1379
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: HWU, PATRICK; NISHIMURA,
APPLICANT: MICHAEL; ROSENBERG, STEVEN A.
TITLE OF INVENTION: T-CELL RECEPTORS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FASTSGY VERSION 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/758,306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               212/132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT ACLIANCE
APPLICATION NUMBER: US/U8/120,JU
FILING DATE: December 3, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: WAIDLIG, RICHARD J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 212/1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1999-1600
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-411-098-35/c
; Sequence 35, Application US/08411098
; Patent No. 5830755
                     879 CAAGIICCAGGAGCIG 894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  625 GACCAGCTCCAGGAGC 640
                                                                     3 cáccicicacidade 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY:
US-08-758-306-971
                                                                                                                                                                      US-08-758-306-971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13;
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                                                                                                                                                                                                                                                                                                                                                                              2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: ALCOUNTRY: United States
COUNTRY: United States
ZIN: 22313-1404
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/990,818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.9%; Score 12.8; DB 1; Best Local Similarity 87.5%; Pred. No. 1.8e+02; Matches 14; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                      0.9%; Score 12.8; DB 1;
87.5%; Pred. No. 1.8e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 40. Application US/08990818

Sequence 40. Application US/08990818

Bartont No. 5910432

GENERAL INFORMATION:

APPLICANT: TYMAKI, Toshitumi

APPLICANT: YAMAKI, Toshitumi

APPLICANT: TSHUOKA, Miyuki

STREET: P.O. BOX 1404

CITY: Alexandria
                                                                                                                                            ; TOPOLOGY: linear MLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "synthetic DNA" (15-08-990-818-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "synthetic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PELLING DATE:
CLASSIFCATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/800,751
FILING DATE:
APPLICATION NUMBER: 08/800,751
FILING DATE: 14-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: TESKIN, RObin L.
REGISTRATION NUMBER: 35,03
REFERENCE/DOCKET NUMBER: 028022-
TELEPHONE: (703) 836-2021
TELEPHONE: (703) 836-2021
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
LEMETH 18 Dase pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                        879 CAAGTTCCAGGAGCTG 894
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 CACGATCCAGGAGCTG 18
                                                                                                                                                                                                                                                                                                                                                                          14; Conservative
                 SEQUENCE CHARACTERISTICS:
                                            LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 233
US-08-990-818-40
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                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/990,818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 39, Application US/08990818
Patent No. 5910432
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: TO, Kiyoshi
APPLICANT: ARIL' Teruo
APPLICANT: ARIL' Teruo
APPLICANT: ARIL' Teruo
APPLICANT: NAKAWURA, Takeshi
TITLE OF INVENTION: NOVEL NITRILE HYDRATASE
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSER: BURNS, DOANE, SWECKER & MATHIS
STREET: P.O. Box 1404
CITY: Alexandria
STRATE: Virginia
COMPUTER: United States
COMPUTER: READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IRM PC COMPATIBLE
COMPATIBLE
COMPUTER: IRM PC COMPATIBLE
COMPUTER: IR
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CLASSIFICATION DATA:
APPLICATION NUMBER: 08/800,751
APPLICATION NUMBER: 08/800,751
APPLICATION NUMBER: 08/800,751
APPLICATION NUMBER: 08/800,751
APPLICATION NUMBER: 07/8010
RAME: TESKIN, RObin L.
REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 35,030
TELECOMMUNICATION:
                                                                                                                                     29,959
ER: 4615-0042.00
      US 08/279,980
                                                                                     ALORNEL MURABHIGE, KATE H.

REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 4615
REFERENCE/DOCKET NUMBER: 4615
TELEPHONE: (202) 887-1500
TELEPHONE: (202) 887-1500
TELERX: (202) 887-0763
TELEX: 90-4030
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          368 TGGGGCCCAGCTTCC 383
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TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: 39:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 TIGGGGCCCAGCGICC 18
                              FILING DATE: 22-JUL-1994
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IMMEDIATE SOURCE:
CLONE: AB3781
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 232
US-08-990-818-39
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Diagnosis, Therapy and Cellular and
Animal Models for Diseases Associated With Mitochondrial
Defects
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0
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Pred. No. 1.8e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/413,740A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Kenyon & Kenyon
STREET: 1025 Connecticut Avenue, N.W.
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/04063
FILING DATE: 30-MAR-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: BORDARY, DAVIG B.
REGISTRATION NUMBER: 34297
REFERENCE/DOCKET NUMBER: 2105/7
FELECOMMUNICATION INFORMATION:
METEROMOTION INFORMATION:
METEROMOTION INFORMATION:
METEROMOTION INFORMATION:
METEROMOTION INFORMATION:
METEROMOTION INFORMATION:
METEROMOTION INFORMATION:
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ZIP: 20036-5405
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Ploppy disk
MEDIUM T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 236
US-08-413-740A-28
Sequence 28, Application US/08413740A
Patent No. 6171859
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MOLECULE TYPE: other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: HERRNSTADT, CORINNA
APPLICANT: PARKER, WILLIAM D.
APPLICANT: DAVIS, ROBERT
APPLICANT: MILLER, SCOTT W.
                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   368 TGGGGGCCCAGCTTCC 383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 TTGGGGCCCAGCGTCC 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (202) 429-177
TELEFAX: (202) 429-0796
INFORMATION FOR SEQ ID NO: 2
                                   (202) 887-0763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.99
Best Local Similarity 87.57
Matches 14; Conservative
                                                                             TELEX: 90-4030
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: DIAGRAM TITLE OF INVENTION: Anim
TITLE OF INVENTION: Defended of the control of the control
                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HYPOLHES NO ANTI-SENSE: NO IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; CLONE: AL
US-09-189-583-18
                       TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   à
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Sequence 34, Application US/09205144
Sequence 34, Application US/09205144
Sequence 34, Application US/09205144
Sequence 34, Application US/09205144
GENERAL INFORMATION:
APPLICANT: Elizabeth J. Ackermann
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF CELLULAR INHIBITOR OF APOPTOSIS-2 EXPRESS
CURRENT APPLICATION NUMBER: US/09/205,144
CURRENT PILING DATE: 1998-12-03
NUMBER OF SEQ ID NOS: 47
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18, Application US/09189583
Patent No. 6063614
GENERAL INFORMATION:
APPLICANT: SELTEN, GERARDUS CORNELIUS MAIA
APPLICANT: SWINKELS, BART WILLEM
APPLICANT: VAN GORCOM, ROBERTUS
APPLICANT: VAN GORCOM, ROBERTUS
APPLICANT: SWINKELS, BARTA
TITLE OF INVENTION: SELECTION MARKER GENE FREE RECOMBINANT
TITLE OF INVENTION: STRAINS: A METHOD FOR OBTAINING THEM AND THE USE OF THESE
TITLES OF INVENTION: STRAINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 37

CORRESPONDENCE ADDRESS:
ADDRESSE: MORRISON & FORRSTER
STREET: 2000 Pennsylvania Ave. N.W., Suite 5500
CITY: Washington, D.C.
COUNTRY: USA
ZIP: 20006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMpatible
OOMPUTER: IBM PC COMPATIBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/189,583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/880,557
FILING DATE: 23-JUN-1997
APPLICATION NUMBER: US 08/279,980
FILING DATE: 22-JUN-1994
ATTORNEY/AGENT INFORMATION:
NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 4615-0042.00
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        257 ACCICCIGGGCIGGCI 272
3 CACGGTCCAGGAGCTG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 Arcrcridecrerci 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; UTHER INFORM
US-09-205-144-34
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FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY, ACENT INFORMATION:
NAME: Warburg, Richard J.
REGIETRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 955-0440
TELERAK: (213) 955-0440
           APPLICATION NUMBER: US/08/584,040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 239
12.08-584-040-8378
Sequence 8378, Application US/08584040
Patent No. 6346598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   218/064
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 GGAGCCAGCUGCUUUU 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 56.21
Best Local 9, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-584-040-3044
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                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                             Sequence 85. Application US/09474922A
Patent No. 6187586
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowsert
APPLICANT: Richard A. Roth
TITLE OF INVENTION:
RILE REPERENCE:
RIS-0036
CURRENT APPLICATION NUMBER: US/09/474,922A
CURRENT FILING DATE: 1999-12-29
NUMBER OF SEQ ID NOS: 89
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                                                                                                 Query Match 0.9%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 1.8e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.9%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 1.8e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 3044, Application US/08584040

Patent No. 6346398
GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: Stinchcomb, Dan T.
APPLICANT: STILLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon
STREET: Sulte 4700
STREET: Sulte 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Antisense Oligonucleotide US-09-474-922A-85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage COMPUTER: IBM Comparible OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                              590 TGCCCCCCACCAGCCT 605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      375 CCAGCTTCCTCCAGAG 390
                                                                                                                                                                                                                                               1 TGCCGCCACCATCCT 16
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COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         U.S.A.
                           8
HYPOTHETICAL:
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US-08-584-040-3044
                         ; ANTI-SENSE:
US-08-413-740A-28
                                                                                                                                                                                                                                                                                                                                                                  US-09-474-922A-85
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 85
LENGTH: 18
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Gaps
                                                       0;
     Length 18;
                                                       2; Indels
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APPLICANT: Pavco, Pamela
APPLICANT: Bavco, Pamela
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Becobedo, Jaime
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 80.02
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Suite 4700
CITY: Los Angeles
STREET: Suite 4700
CITY: Los Angeles
STREET: Suite 4700
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER REAAABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: John DayAR:
MEDIUM TYPE: John DayAR:
MEDIUM TYPE: John DayAR:
MEDIUM TYPE: John DAYA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION NUMBER: G0/005,974
FILING DATE: Gatchard J.
REPERBRACH OFFERENT MINERER: 3210764
REPERBRACH OFFERENT MINERER: 3210764
0.9%; Score 12.8; DB 1;
56.2%; Pred. No. 1.8e+02;
tive 5; Mismatches 2;
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Gaps
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0.9%; Score 12.8; DB 1; Length 18;
Best Local Similarity 68.8%; Pred. No. 1.8e+02;
Matches 11; Conservative 3; Mismatches 2; Indels
                                                                                   0.9%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 1.8e+02;
                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Zwick, Michael G.
APPLICANT: Zwick, Michael G.
APPLICANT: Edington, Brent E.
APPLICANT: McSwiggen, James A.
APPLICANT: McSwiggen, James A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Young, Scott A.
APPLICANT: Folkerts, Otto
APPLICANT: Folkerts, Otto
APPLICANT: McTlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR
TITLE OF INVENTION: MODULATION OF GENE EXPRESSION
TITLE OF INVENTION: 10 PLANTS
NUMBER OF SEQUENCES: 1263
                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTY: u.o...

ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: LDM Compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/675,645
FILING DATE: July 12, 1996
CLASSIFICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Waxburg Richard J.
REGISTRATION NUMBER: 32,327
TELECOMMUTICATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                          Sequence 629, Application US/08679645; Patent No. 6350934; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                884 TCCAGGAGCTGCGGTA 899
                                                                                                                                                                                                                          16 rccardaccrdcccca 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (213) = TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: SULCE CITY: Los Angeles STATE: California
                                                                              Query Match
Best Local Similarity
Matches 14; Conserv
                      i
US-08-679-645-609
                                                                                                                                                                                                                                                                                                                         US-08-679-645-629
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0
                                                                                                                                                                                                                                        0.9%; Score 12.8; DB 1; Length 18; 56.2%; Pred. No. 1.8e+02; tive 5; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Skokut, Thomas A.
APPLICANT: Young, Scott A.
APPLICANT: Young, Scott A.
APPLICANT: Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR TITLE OF INVENTION: MODULATION OF GENE EXPRESSION TITLE OF INVENTION: IN PLANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A.

ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IRM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
CLASSIFTCATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELEFAX: (213) 955-0440
TELEFAX: 67-3510
INFORMATION FOR SEQ ID NO: 609:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zwick, Michael G.
Bdington, Brent E.
McSwiggen, James A.
Merlo, Patricia Ann Owens
Guo, Lining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 609, Application US/08679645
Patent No. 6350934
GENERAL INFORMATION:
APPLICANT: Zwick, Michael G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Lyon E Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                   1311 GTAGCCAGGTGCTTTT 1326
                                                                                                                                                                                                                                                                                                                                                              | |||||| :||::::
1 GGAGCCAGCUGCUUUU 16
                      TELEX: 67.3510
INFORMATION FOR SEQ ID NO: 8
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1263
       (213) 955-0440
                                                                                                                                                                                                                     LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: LYON & LYON
STREET: 833 West Fiftl
STREET: Suite 4700
                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-679-645-609/c
                                                                                                                                                                      ; TOPOLOGY: 1
US-08-584-040-8378
TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
APPLICANT:
APPLICANT:
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Sequence 7504, Application US/09422978

Sequence 7504, Application US/09422978

BEREARL INFORMATION:
APPLICANT: Cohen, Daniel
APPLICANT: Chumenfeld, Marta
CURRENT PRINTON: Biallelic markers for use in constructing a high density...
FILE REFERENCE: GENSET. 020CP1
CURRENT PAPLICATION NUMBER: US 09/298,850
EARLIER APPLICATION NUMBER: US 09/298,850
EARLIER PILING DATE: 1999-04-21
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER PILING DATE: 1998-11-23
                                  APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF CYCLIN D2 EXPRESSION
FILE REFERENCE: RTS-0275
CURRENT APPLICATION UNBER: US/09/920,760
CURRENT FILING DATE: 2001-08-01
NUMBER OF SEQ ID NOS: 89
LENGIH: 18
                                                                                                                                                                                                                                                                                                                                                                                                Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17, Application US/09077619
Patent No. 6500614
GENERAL INFORMATION:
APPLICANT: ARGUELLO, Rafael
APPLICANT: AVAKTAN, Hovanes
APPLICANT: ANAKTAN, Alejandro
1TILE OF INVENTION: METHOD FOR IDENTIFYING AN UNKNOWN ALLELE
FILE REFERENCE: 028979/0104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.9%; Score 12.8; DB 1; Best Local Similarity 87.5%; Pred. No. 1.8e+02; Matches 14; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                   ; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-920-760-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/077,619
CURRENT FILING DATE: 2000-03-31
PRIOR APPLICATION NUMBER: PCT/GB96/02959
PRIOR FILING DATE: 1996-11-29
PRIOR FILING DATE: 1995-11-29
NUMBER OF SEQ ID NOS: 46
SCOTTWARE: Patentin version 3.0
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1145 TITITICITITIGGAA 1160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           531 GGAGCAGCTGGGTGCC 546
                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 rrrrgrrrrrrddaa 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GGAGCAGCTGAGAGCC 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-422-978-7504
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US-09-077-619-17
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                                                                                                                                                                    Sequence 51, Application US/08614151
Parent No. 6436635
Parent No. 6436635
Parent No. 6436635
APPLICANT: CANTOR, Charles R.
APPLICANT: KOSTER, Hubert
APPLICANT: SMITH, Cassandra L.
TITLE OF INVENTION: SOLID PHASE SEQUENCING OF DOUBLE-STRANDED NUCLEIC ACID
NUMBER OF SEQUENCES: 53
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ..
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87.5%; Pred. No. 1.8e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                     STREET: 1299 Pennsylvania Avenue, N.W. CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 20004-2400
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: EDM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FESTEN: DOS
SOFTWARE: FASTEN: US/OS/614,151
FILING DATE: 12-MAR-1996
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/420,009
FILING DATE: 11-APR-1995
APPLICATION NUMBER: 08/110,691
FILING DATE: 23-AUG-1993
APPLICATION NUMBER: 08/110,691
FILING DATE: 23-AUG-1993
APPLICATION NUMBER: 08/110,691
FILING DATE: 3-AUG-1993
APPLICATION NUMBER: 07/972,012
FILING DATE: 36-NOV-199
APPLICATION NUMBER: 06-NOV-199
AFICHER REMEMBER: 36,902
REFERENCE/DOCKET NUMBER: 36,902
REFERENCE/DOCKET NUMBER: 36,902
REFERENCE/DOCKET NUMBER: 36,902
REFERENCE/DOCKET NUMBER: 16865-0276
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
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US-09-920-760-14
Sequence 14, Application US/09920760
; Parent No. 6492173
220 CGAGCTCCTCAGCCTC 235
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                             CGUGCUGCUCAGCCUC 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.98
Best Local Similarity 87.59
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 202-639-7890
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MOLECULE TYPE: CDNA
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ORIGINAL SOURCE:
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Gaps

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APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PILING DATE: 1956-10-26
PRIOR FILING DATE: 1956-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Parco, Pam
APPLICANT: Stinchcomb, Dan
APPLICANT: Scinchcomb, Dan
APPLICANT: Scinchcomb, Dan
APPLICANT: Stinchcomb, Daime
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Scooled, Jaime
APPLICANT: Scooled, Jaime
APPLICANT: Scooled, Jaime
APPLICANT: Scooled, Jaime
APPLICANT: Stinchcomb, Dan
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APPLICANT: Scooled, Jaime
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcom
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OTHER INFORMATION: Description of Unknown Organism: Primer,
OTHER INFORMATION: genomic DNA
US-09-742-313-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.9%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred. No. 1.8e+02;
Matches 14; Conservative 0; Mismatches 2;
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56.2%; Pred. No. 1.8e+02;
ive 5; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1472, Application US/09371772B
Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
           PRIOR APPLICATION NUMBER: 10016278.9
PRIOR FILING DATE: 2000-04-03
PRIOR APPLICATION NUMBER: 10027954.6
PRIOR FILING DATE: 2000-06-08
NUMBER OF SEQ ID NOS: 12
SOGTWARE: Patentin Ver. 2.1
SEQ ID NO 6
LENGTH: 18
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GGAGCCAGCUGCUUUU 16
                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Unknown Organism
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US-09-371-772B-1472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 9; Conserv
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US-09-422-978-11175

Sequence 11075, Application US/09422978

Patent No. 6537751

GENERAL INFORMATION:

APPLICANT: Cohen, Daniel

APPLICANT: Chumakov, Ilya

TITLE OF INVENTION: Biallelic markers for use in constructing a high density...

FILE REFERENCE: GENSET.020791

CURRENT APPLICATION NUMBER: US/09/422,978

CURRENT APPLICATION NUMBER: US 09/298,850

BARLIER APPLICATION NUMBER: US 60/109,732

BARLIER APPLICATION NUMBER: US 60/109,732

BARLIER APPLICATION NUMBER: US 60/109,732

BARLIER PILING DATE: 1998-04-21

BARLIER PILING DATE: 1998-04-21

BARLIER PILING DATE: 1998-04-21

BARLIER PILING DATE: 1998-04-21

SEALLIER PILING DATE: 1998-04-21

SEALLIER PILING DATE: 1998-04-21

SEQ ID NO 11175

LENGTH: 18
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. OTHER INFORMATION: downstream amplification primer 99-3147 for SEQ 3310, in compleme
US-09-422-978-11175
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Patent No. 6562946
GENERAL INFORMATION:
APPLICANT: Althaus, Harald
APPLICANT: Hauser, Hans-Peter
TITLE OF INVENTION: Human Procalcitonin and the Preparation and Use Thereof
FILE REFERENCE: 0552.1445.00
CURRENT APPLICATION NUMBER: 1090/742,373
CURRENT PILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 19962434.8
PRIOR FILING DATE: 1999-12-22
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                                                                                                                                                                                                                                                                                                                   ; LOCATION: 1..18 ; OTHER INFORMATION: upstream amplification primer 99-6071 for SEQ 3570, US-09-422-978-7504
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
0.9%; Score 12.8; DB 1; Length 18;
Best Local Similarity 87.5%; Pred. No. 1.8e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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EARLIER APPLICATION NUMBER: US 60/082,614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         850 TCAGCATACCGCTTTG 865
                              EARLIER FILING DATE: 1998-04-21
NUMBER OF SEQ ID NOS: 11796
SEQ ID NO 7504
LENGTH: 18
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                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                              NAME/KEY: primer_bind LOCATION: 1.18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: primer_bind
LOCATION: 1..18
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Diagnosis, Therapy and Cellular and
Animal Models for Diseases Associated With Mitochondrial
Defects
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;Patent No. 5182195
; APPLICANT: NAKHHAMA, KAZUO;KAISHO, YOSHIHIKO;YOSHIMURA, KOJI
; TITLE OF INVENTION: METHOD FOR INCREASING USING PROTEASE
;DEPICIENT YEASTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/04063
FILING DATE: 30-MAR-1995
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: BOOMHAM, David B.
REGISTRATION NUMBER: 34.997
REFERENCE/DOCKET NUMBER: 2105/7
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 429-1776
TELEPHONE: (202) 429-0796
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.9%; Score 12.8; DB 1;
87.5%; Pred. No. 1.8e+02;
tive 0; Mismatches 2;
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/269,140
FILING DATE: 09-NOV-1988
                                                                                                                                                        Application PC/TUS9504063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 20036-5405
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                other nucleic acid
                                                                                                                                                   Sequence 28, Application PC/TUS950406
GENERAL INFORMATION:
APPLICANT: HERRNSTADT, CORINNA
APPLICANT: PARKER, WILLIAM D.
APPLICANT: DAVIS, ROBERT
APPLICANT: MILLER, SCOTT W.
ITLE OF INVENTION: Diagnosis, TP
TITLE OF INVENTION: Animal Models
ITLE OF INVENTION: Animal Models
ITLE OF SEQUENCES: 206
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                  Kenyon & Kenyon
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                 2 CCCCACATCCCAGTTC 17
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Matches 14, Conservative
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EDNESS: double
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                                                                                                                                    r-US95-04063-28
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PCT-USS3-12600-16
PCT-USS3-12600-16
SEQUENCE 16, Application PC/TUS9312600
SEQUENCE 16, Application PC/TUS9312600
SEQUENCE 16, Application:
APPLICANT: Dixon, Richard A.F.
APPLICANT: Dixon, Richard A.F.
TITLE OF INVENTION: ANTISENSE MOLECULES DIRECTED AGAINST A
TITLE OF INVENTION: ANTISENSE MOLECULES DIRECTED AGAINST A
TITLE OF INVENTION: ANTISENSE
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSE: Milnamow, Lid.
STREET: 180 North Stetson, Suite 4700
STREET: Chicago
STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                      0.9%; Score 12.8; DB 1; Length 18; 56.2%; Pred. No. 1.8e+02; tive 5; Mismatches 2; Indels
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SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/12600
FILING DATE: 28-DSC-1993
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/999,706
FILING DATE: December 31, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Katz, Martin L.
REGISTRATION NUMBER: 25,011
TELEPHONE: (312/616-5400
TELEPHONE: (312/616-5400
TELEPHONE: (312/616-5400
TELEPHONE: (312/616-5400
TELEPHONE: (312/616-5400
TELEPHONE: (312/616-5400
TELEPHONE: US SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
CURRENT APPLICATION NUMBER: US/09/311,772B CURRENT FILING DATE: 1999-08-10 PRIOR APPLICATION NUMBER: US 60/005,974 PRIOR FILING DATE: 1995-10-26 PRIOR FILING DATE: 1996-10-08 PRIOR FILING DATE: 1996-01-08 NUMBER OF SEQ ID NOS: 14225 SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 60601
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                   1311 GTAGCCAGGTGCTTTT 1326
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Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
                                                                                                                                                                                                                                           TYPE: RNA
ORGANISM: Mus sp.
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PCT-US93-12600-16
                                                                                                                                                                                                                                                                                       US-09-371-772B-4034
                                                                                                                                                                                          SEQ ID NO 4034
LENGTH: 18
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LENGTH: 18

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870 CCCCACAGCCAAGTIC 885

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Conservative

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US-09-661753-55/c

| Sequence 55, Application US/09661753 |
| Patent No. 6436909 |
| Patent No. 6436909 |
| Patent No. 6436909 |
| APPLICANT: Susan R. Muzray |
| APPLICANT: Susan R. Muzray |
| TITLE OF INVENTION: ANTISENSE MODULATION OF TRANSFORMING GROWTH FACTOR BETA |
| PILE REFERENCE: ISPH-0498 |
| CURRENT APPLICATION NUMBER: US/09/661,753 |
| CURRENT APPLICATION NUMBER: 60/154,546 |
| FARLIER APPLICATION NUMBER: 60/154,546 |
| FARLIER OF SEQ ID NOS: 68 |
| SEQ ID NO 55 |
| LENGTH: 20 |
| LENGTH: 20 |
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APPLICANT: Pardinas, J.
APPLICANT: Pardinas, J.
APPLICANT: Partimoo, S.
APPLICANT: Prouty, S.
APPLICANT: STORY S.
APPLICANT: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE: OTHER INFORMATION: Description of Artificial Sequence: primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 12.6; DB 1; Length 20;
Pred. No. 2.5e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.9%; Score 12.4; DB 1; Length 14; 92.9%; Pred. No. 1.2e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FRATURE:

// OTHER INFORMATION: Antisense Oligonucleotide
US-09-661-753-55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-832-021-15
; Sequence 15, Application US/08832021
; Patent No. 6045998
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Best Local Similarity 78.9%;
Matches 15; Conservative C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1145 TITITICITITIGG 1158
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Best Local Similarity 92.99
Matches 13; Conservative
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LENGTH: 14
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                  Length 18;
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                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mateuraki, Hajime
Macall, Glenn H.
INVENTION: NUCLEIC ACID ANALYSIS TECHNIQUES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDUW TYPE: FLORM:
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Joe Liebeschuetz
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/882,649A
FILING DATE: 25-0un-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/000
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/010,471
FILING DATE: 23-0AN-1996
APPLICATION NUMBER: US 60/035,170
FILING DATE: 09-0AN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Liebeschuetz, Joe
REGISTRATION NUMBER: 37,505
REGISTRATION NUMBER: 37,505
REGISTRATION NUMBER: 37,505
REGISTRATION NUMBER: 37,505
0.9%; Score 12.8; DB 1;
87.5%; Pred. No. 1.8e+02;
tive 0; Mismatches 2;
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; SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-08-882-649A-7
                                                                                                                                                                                                                                                                                                                                                   RESULT 253
US-08-882-649A-7/c
Sequence 7, Application US/08882649A
Setent No. 6344316
GENERAL INFORMATION:
CHERNORMATION:
GHERAL TOOKHART, David J.
CHERNORMATION:
CHERNORM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wodicka, Lisa
Cronin, Maureen T.
Lee, Danny
Tran, Huu M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                       473 AGGGGGAGGACTGCCG 488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: San Francisco
STATE: CA
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                                                                                     14; Conservative
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Best Local Similarity 86.7.
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
                                        Best Local Similarity
Matches 14; Conserva
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    Query Match
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US-08-985-162-1842; Sequence 1842, Application US/08985162; Patent No. 6057156; GENERAL INFORMATION:

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Gaps

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1144 TTTTTTTTTTGG 1158

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Sequence 12, Application US/08882164D

Patent No. 6306624

GENERAL INFORMATION:

APPLICANT: Beckett, Barbara R., Jones, Glenville

TITLE OF INVENTION: Retinoid Metabolizing Protein

NUMBER OF SEQUENCES: 43

CORRESPONDENCE ADDRESS:

ADDRESSEE: Blake, Cassels & Graydon

STREET: Dox 25, Commerce Court West

CITY: Toronto

COUNTRY: Canada

ZIP: MEILING

STATE: Ontario

COMPUTER: Canada

ZIP: MEILING

COMPUTER: CANBADLE FORM:

MEDIUM TYPE: Diskette, 3 1/2 inch, 1.4 Mb storage

COMPUTER: COMPAQ, IBM PC compatible

COMPUTER: ONDEACH IN MS-DOS 5.1

SOFTWARE: WORD PERRECT

CURRENT APPLICATION DATE:

APPLICATION NUMBER: US/08/882,164D

FILING DATE: June 25, 1997

PRIOR APPLICATION DATE:

APPLICATION NUMBER: 08/667,546

FILING DATE: October 1, 1996

APPLICATION NUMBER: 06/6724,466

FILING DATE: October 1, 1996

APPLICATION NUMBER: 07/24,466

FILING DATE: October 1, 1996

APPLICATION NUMBER: 07/24,466

FILING DATE: OCTOBER 1, 1996

APPLICATION NUMBER: 07/24,466

FILING DATE: OCTOBER 1, 1996

APPLICATION NUMBER: 56,424

REGISTRATION NUMBER: 56,424

REFERENCE/DOCKET NUMBER: 56,727/00000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.9%; Score 12.4; DB 1; Best Local Similarity 92.9%; Pred. No. 1.2e+02; Matches 13; Conservative 0; Mismatches 1;
                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/667,546
FILING DATE: Unne 21, 1996
ATTORNEY AGENT INFORMATION:
NAME: Hunt, John C.
REGISTRATION NUMBER: 36,424
REPERENCE/DOCKET NUMBER: 50767/00004
TELECOMMUNICATION INFORMATION:
TELECHONE: (416) 863-4344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 50767/00010
TELECOMMUNICATION INFORMATION:
                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/724,466B
FILING DATE: October 1, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1145 TITITICITITIGG 1158
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TELEFAX: (416) 863-2653
INFORMATION FOR SEQ ID NO: 12
SEQUENCE CHARACTERISTICS:
LENGTH: 14 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (416) 863-4344
                                                                                                                                                                                                                                                                                                                             TELEFAX: (416) 863-2653
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 14 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TOPOLOGY: linear
US-08-724-466B-12
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US-08-882-164D-12
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Akhtar, Saghir
Fell, Patricia
McSwiggen, James
VENTION: ENZYMATIC NUCLEIC ACID TREATMENT
VENTION: OF DISEASES OR CONDITIONS RELATED
VENTION: PACTOR RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 12, Application US/08724466B
Parent No. 6063606
GENERAL INFORMATION:
APPLICANT: Petkovich, P. Martin, White, Jay A.,
APPLICANT: Beckett, Barbara R., Jones, Glenville
TITLE OF INVENTION: Rectinoid Metabolizing Protein
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE Blake, Cassels & Graydon
STREDT: Box 25, Commerce Court West
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: TOLUN.

ZIP: M5L 1A9
ZONTRY: Canada
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3 1/2 inch, 1.4 Mb storage
COMPUTER: COMPAQ, IBM PC compatible
COMPUTER: CAPEM: MS-DOS 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.9%; Score 12.4; DB 1;
85.7%; Pred. No. 1.2e+02;
ttive 1; Mismatches 1;
                                                             TITLE OF INVENTION: ENZYMATIC NUCLEIC ACTIVE OF INVENTION: OF DISEASES OR CONDITIONS OF DISEASES OR CONDITIONS OF DISEASES OR CONDITIONS OF DISEASES OR CONDITIONS OF ENTRE OF ENVENTION: TO LEVELS OF EPIDERM VIMBER OF SEQUENCES: 1877

CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STREET: U.S.A.
ZIP: 90701-2066
COMPUTER REDABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 MD
                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IDBM P.C. DOS 5.0
SOFTWARE: FastSEQ for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/985,162
FILING DATE: 04 December 1997
CLASSIFICATION: 514
PRICR APPLICATION: 514
PRICR APPLICATION: DATA:
APPLICATION NUMBER: 60/05,476
FILING DATE: 31 January 1997
APPLICATION NUMBER: 32,327
ATTORNEY/AGENT INPORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 230/107
TELECOMMUNICATION INFORMATION:
TELECHOME (213) 955-0440
TELEPHONE: (213) 955-0440
TELEPHONE: 67-3510
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1050 CGACAGCCCTGGCC 1063
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Best Local Similarity 85.7%
...hes 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-985-162-1842
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20,532
ER: 11859-1
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    RESULT 260
US-08-863-639A-7/c
; Sequence 7, Application US/08663639A
; Patent No. 5991185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY AGENT 1875
ATTORNEY AGENT 1875
NAME: JOSEPh E Mueth
REGISTRATION NUMBER: 20,532
REFERENCE/DOCKET NUMBER: 1185:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (626) 796-400
TELEPHONE: (626) 795-631
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1144 TTTTTTTTTTG 1157
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Best Local Similarity 92.99
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 91101
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single
                                                                                                                                                                                                                                                                                                                                     Pasadena
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US-08-832-021-50
                                                                                                                                                                                                                                                                                                                                                           STATE:
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                                         Query Match 0.9%; Score 12.4; DB 1; Length 14; Best Local Similarity 92.9%; Pred. No. 1.2e+02; Matches 13; Conservative 0; Mismatches 1; Indels
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0.9%; Score 12.4; DB 1; Length 15;
Best Local Similarity 92.9%; Pred. No. 1.4e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                    WESONE 23, Application US/08319492B

Sequence 23, Application US/08319492B

Patent No. 561648B

GENERAL INFORMATION:
APPLICANT: Sullivan, Sean M.
APPLICANT: Draper, Kenneth G.
APPLICANT: Stinchcomb, Dan T.
TITLE OF INVENTION: RIBOZYME TREATMENT OF DISEASES
TITLE OF INVENTION: OF COUDITIONS RELATED TO LEVELS
TITLE OF SEQUENCES: 751

NUMBER OF SEQUENCES: 751

NUMBER OF SEGUENCES: 751

STREET: 633 West Fifth Street
STREET: Suite 4700

CITY: LOS ANGELSE

START: LOS ANGELSE

STREET: Suite 4700

CITY: LOS ANGELSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/319,492B
FILING DATE: October 7, 1994
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: including application
APPLICATION DATA: including application
APPLICATION DATA: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 09/098,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PEFFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REPRENCE/DOCKET NUMBER: 209/.
TELECOMMUNICATION INPORMATION:
TELEPHONE: (213) 955-0440
TELETAX: (213) 955-0440
TELETAX: (7-3510
INPORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                        1145 TTTTTTCTTTTGG 1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   628 CAGCTCCAGGAGCT 641
                                                                                                                                                                                      1 TITITITITITIGG 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      U.S.A.
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US-08-882-164D-12
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STATE: Ca
COUNTRY:
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Gaps
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| Sequence 50, Application US/08832021
| Patent No. 604598
| GENERAL INFORMATION:
| APPLICANT: Combates, J.
| APPLICANT: Partimoo, S.
| APPLICANT: Prouty, S.
| APPLICANT: Stenn, K.
| TITLE OF INVENTION: IMPROVED TECHNIQUE FOR DIFFERENTIAL DISPLAY
| FILE REFRENCE: JBP-382
| CURRENT FILING DATE: 1997-04-02
| SOUTRARY FILING DATE: 1997-04-02
| SOUTRARE PATENTING DATE: 1997-04-02
| SOUTRARE PATENTIAL OF SEQ ID NOS: 64
| SOUTRARE PATENTIAL OF SEQ ID NOS: 64
| SEQ ID NO 50
| LENGTH: 15
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GENERAL INFORMATION:
APPLICANT: Matson, Robert S.
APPLICANT: Cassin, Peter J.
APPLICANT: Rampal, Jang B.
APPLICANT: Caskey, C. T.
TITLE OF INVENTION: OLIGONUCLEOTIDE REPEAT ARRAYS
NUMBER OF SEQUENCES: 95
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sheldon & Mak
STREET: 225 South Lake Avenue, 9th Floor
                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: Windows 95 SOFTWARE: Corel WordPerfect 8 version CURRENT APPLICATION DATA: US/08/863,639A FILLNG DATE: My 28, 1997 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.9%; Score 12.4; DB 1;
92.9%; Pred. No. 1.4e+02;
tive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence
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Length 15;
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                                                                                                                                                                                                                                  APPLICANT: Kiely, John
APPLICANT: Kiely, John
APPLICANT: Griffin, Michael
APPLICANT: Griffin, Michael
APPLICANT: Griffin, Michael
APPLICANT: Griffin, Michael
APPLICANT: Buchardt, Ole
APPLICANT: Christensen, Leif
ITILE OF INVENTION: Linked Peptide Nucleic Acids
FILE REFERENCE: 18131577
CURRENT APPLICATION NUMBER: 08/08/25,951
PRIOR APPLICATION NUMBER: 08/08,658
PRIOR PILING DATE: 1993-07-02
PRIOR PILING DATE: 1991-05-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.9%; Score 12.4; DB 1;
92.9%; Pred. No. 1.4e+02;
tive 0; Mismatches 1;
                                                                                                                                                                                                        ; Sequence 31, Application US/08275951; Patent No. 6451968; GENERAL INFORMATION:
        1145 TTTTTTTTTGG 1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1142 CCTTTTTTTTTT 1155
                                                              1 rrrrrrrrrrrss 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                               RESULT 264
US-08-275-951-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 265
US-08-087-387-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                             Sequence 5. Application US/08832021
; Sequence 5. Application US/08832021
; Patent No. 604598
; GENERAL INFORMATION:
    APPLICANT: Combates, N.
    APPLICANT: Pardinos, J.
    APPLICANT: Pardinos, S.
    APPLICANT: Patent, S.
    APPLICANT: Patent, S.
    APPLICANT: Patent, S.
    APPLICANT: Patent, S.
    APPLICANTON: IMPROVED TECHNIQUE FOR DIFFERENTIAL DISPLAY
    TILE REFERENCE: JBP-382
    CURRENT APPLICATION NUMBER: US/08/832,021
    CURRENT FILING DATE: 1997-04-02
    NUMBER OF SEQ ID NOS: 64
    SOFTHARE: Patent In Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Description of Artificial Sequence: primer US-08-832-021-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: primer
US-08-832-021-51
; OTHER INFORMATION: Description of Artificial Sequence: primer US-08-832-021-50
                                                                                Query Match 0.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 1.4e+02; Matches 13; Conservative 0; Mismatches 1; Indels
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Query Match

Best Local Similarity 92.9%; Pred. No. 1.4e+02;
Matches 13; Conservative 0; Mismatches 1; Indele
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0.9%; Score 12.4; DB 1; Length 15;
Best Local Similarity 92.9%; Pred. No. 1.4e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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Patent No. 6045998
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
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LENGTH: 15
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Sequence 6, Application US/08087387

Patent No. 5473060

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Sergei M. Gryaznov
ITILE OF INVENTION:
ORRESPONDENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Stephen C. Macevicz, Lynx Therapeutics
STREET: 465 Lincoln Centre Drive
STREET: Foster City
STREET: California
) OTHER INFORMATION: Description of Artificial Sequence: No. 6451968el Sequence: NAME/KEY: misc_feature
| NAME/KEY: misc_feature
| LOCATION: (6)...(7)
| OTHER INFORMATION: Lysine, Amino Hexanoic Acid, Lysine, Amino OTHER INFORMATION: Hexanoic Acid, Lysine Linkage
| US-08-275-951-31
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                                                                                                                                                                                                                                                                                                                                                                                   US-08-131-365B-23/C

US-08-131-365B-23/C

Sequence 23, Application US/08131365B

Patent No. 5257690

GENERAL INFORMATION:
APPLICANT: Briggs, Michael S.
APPLICANT: Briggs, Michael R.
APPLICANT: Wang, Xiaodong,
APPLICANT: Wang, Xiaodong,
APPLICANT: Goldstein, Joseph L.
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING
TITLE OF INVENTION: TO STEROL REGULATORY ELEMENT BINDING
TITLE OF INVENTION: PROTEINS

ITLLE OF INVENTION: PROTEINS

NUMBER OF SEQUENCES: 64
                                                                                                                                                                                                 Indels
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COUNTRY: U.S.A.

ID: 77210

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/131,365B
FILING DATE: 01-OCT-1993
CLLASSIFICATION: A355
ATTORNEY/AGENT INFORMATION:
NAME: Parker, David L.
REGISTRATION NUMBER: UTSD:372/PAR
TELECOMMUNICATION INFORMATION:
TELEBENOR: (512) 474-7577
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: GOVERNEY lings
                                                                                                                                           Query Match

0.9%; Score 12.4; DB 1;
Best Local Similarity 92.9%; Pred. No. 1.7e+02;
Matches 13; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee
STREET: P.O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 6, Application US/08455627
                                                                ; MOLECULE TYPE: DNA (genomic) US-08-061-697-23
                                                                                                                                                                                                                                             471 GCAGGGGGACT 484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      471 GCAGGGGGAGGACT 484
                                                                                                                                                                                                                                                                                              16 GCAGGGGGAGGT 3
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GCAGGGGGAGGAGT
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Best Local Similarity
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US-08-455-627-6
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Parent No. 5498696

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Xiaodong; Goldstein, Joseph L.
TITLE OF INVENTION: Sterol Regulatory Element Binding Proteins TITLE OF INVENTION: and Their Use in Screening Assays
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEN: ALORESSEN:
TITLE OF INVENTION: MALE & Durkee
CORRESPONDENCE ADDRESS:
ADDRESSEN: ALORESSEN:
CITY: Houston
CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.9%; Score 12.4; DB 1; Length 16; 92.9%; Pred. No. 1.7e+02; tive 0; Mismatches 1; Indels
       COMPUTER READABLE FORM:

MEDIUM TYPE: 5.25 inch diskette

COMPUTER: IBM compactible

OPERATING SYSTEM: Windows 3.1/DOS 5.0

CURRENT APPLICATION DATA:

APPLICATION UNMBER: U$/08/087,387

FILING DATE: 19930702

CLASSIFICATION 435

FRICK APPLICATION DATA:

APPLICATION NUMBER: BATCH APPLICATION BATCH APPLICATION WINDER: BATCH APPLICATION WINDER: BATCH APPLICATION WINDER: BATCH APPLICATION NUMBER: BATCH APPLICATION NUMBER: BATCH APPLICATION NUMBER: BATCH APPLICATION NUMBER: 30,285

REGERRANCE DOCKET NUMBER: 104

TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: BIM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 05/08/061,697
FILING DATE: CONCURRENTLY Herewith
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Parker, David L.
REGISTRATION NUMBER: 32,165
REFERENCE/POCKET NUMBER: 32,165
REFERENCE/COKET NUMBER: 1TELEPHONE: (512) 320-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1143 CTTTTTTTTTT 1156
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (415) 358-7794 INFORMATION FOR SEQ ID NO: (SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 16 nucleotides
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: Tex.
COUNTRY: USA
77210
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US-08-061-697-23/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-087-387-6
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Sequence 6, Application US/08461271
Patent No. 5741643
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Sergei M. Gryaznov
TITLE OF INVENTION: Oligonucleotide clamps having diagnostic
TITLE OF INVENTION: and therapeutic applications
                                                                                                                                                                                                                                                                                                                         Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.9%; Score 12.4; DB 1; Length 16; 92.9%; Pred. No. 1.7e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                    i; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: Stephen C. Macevicz, Lynx Therapeutics STREET: 465 Lincoln Centre Drive
CITY: Foster City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 94404

COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch diskette
COMPUTER: ISM compatible
OPERATING SYSTEM: Windows 3.1/DOS 5.0
SOFTWARE: Microsoft Word for Windows, vers. 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,271
                                                                                                                                                                                                                                                                                                                     Query Match 0.9%; Score 12.4; DB 1;
Best Local Similarity 92.9%; Pred. No. 1.7e+02;
Matches 13; Conservative 0; Mismatches 1;
                            38,534
ER: MSWA-0001
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                                               REFERENCE/DOCKET NUMBER: MSI
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEFAX: 215-568-3439
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Stephen C. Macevicz
REGISTRATION NUMBER: 30,285
REFERENCE/DOCKET NUMBER: 10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 358-7855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/087
FILING DATE: 2-0ul-93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                 1145 TTTTTTTTGG 1158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (415) 358-7794
INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 16 nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                            14 TTTTTTTTTGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13; Conservative
                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
                            REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: California
COUNTRY: USA
                                                                                                                                                                                       LENGTH: 16 bases
                                                                                                                                                                                                                                                       TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                                          US-08-284-484A-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 270
US-08-461-271-6
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Sequence 4, Application US/08284484A
Fatent No. 5639873
GENERAL INFORMATION:
APPLICANT: Barascut, et al.
TITLE OF INVENTION: Oligothionucleotides
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSE: Woodcock Washburn Kurtz Mackiewicz and No. 5639873ris
STREET: One Liberty Place - 46th Floor
CITY: Philadelphia
                                                                                          Synthesis of Branched and Multiply
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
; Patent No. 55716,,
; GENERAL INFORMATION:
; APPLICANT: Sergei M. Gryaznov
; TITLE OF INVENTION: Convergent Synthesis of Brancheu ....
TITLE OF INVENTION: Connected Macromolecular Structures;
NUMBER OF SEQUENCES: 26
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Cooley Godward LLP
GTREET: Five Palo Alto Square, 3000 El Camino Real
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                               STATE:
COUNTRY: USA

ZIP: 94306-215

ZIP: 94306-215

ZIP: 94306-215

ZIP: 94306-215

ZIP: PADELE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/455,627
FILING DATE: 31-MAX-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: NAKAMURA: 35,966
REFERRICE/DOCKET NUMBER: 15,966
REFERRICE/DOCKET NUMBER: 15,966
TELEPHONE: 415-843-5000
"TELEPHONE: 415-843-5000
"TELEPHONE: 415-857-0663
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92.9%; Pred. No. 1.7e+02;
tive 0; Mismatches 1;
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/284,484A
FILING DATE: 04-AUG-1994
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Paul K. Legaard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A.

ZIP: 19103

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 720 Kb
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1143 CTTTTTTTTTTT 1156
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.99
Best Local Similarity 92.99
Matches 13; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: DNA
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Gaps

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COMPUTER: //210
COMPUTER: FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: EN PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/668,123
FILING DATE: 14-JUN-1996
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/131,365
FRICK ASPICATION NUMBER: 08/131,365
FRILNG DATE: 01-OCT-1993
ATTORNEY/AGENT INFORMATION:
NAME: PARKER: DAVIG L.
REGISTRATION NUMBER: 32,165
REFRENCE/DOCKET NUMBER: 32,165
REFRENCE/DOCKET NUMBER: UTSD:372/PAR
TELECOMMULCATION INFORMATION:
TELECOMMULCATION INFORMATION:
TELECOMMULCATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Arnold, White & Durkee STREET: P.O. Box 4433 CITY: Houston CITY: Houston COUNTE: Texas COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 23, Application US/08668123
Patent No. 5891631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.9%;
Best Local Similarity 92.9%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1143 CTTTTTTTTTT 1156
                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 16 nucleotide
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 273
US-08-668-123-23/c
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Patent No. 5830658
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION:
ORRESPONDENCE ADDRESS:
ADDRESSE:
CORRESPONDENCE ADDRESS:
ADDRESSE:
TYRET: Five Palo Alto Square, 3000 El Camino Real
CITY: Palo Alto
STATE: California
  Sequence 6, Application US/08713685A
; Sequence 6, Application US/08713685A
; Patent No. 5817795
; GENERAL INFORMATION:
; APPLICANT: Sergel M. Gryaznov
TITLE OF INVENTION: and therapeutic applications
; TITLE OF INVENTION: and therapeutic applications
; NUMBER OF SEQUENCES: 6
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Stephen C. Macevicz, Lynx Therapeutics
; STREET: 465 Lincoln Centre Drive
CITY: Foster City
; STATE: California
; COUNTRY: USA
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                                                                                                                                                                                                                                                                                                              ZIP: 94404

ZIP: 94404

CMEDTUM TYPER 5.25 inch diskette
COMPUTER: 18M compatible
COMPUTER: INCOMEDIA
OPERATING SYSTEM: Windows 3.1/DOS 5.0
SOFTWARE: Microsoft Word for Windows, vers. 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/713,685A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
CLASSIFICATION: 435
RIOR APPLICATION DATA:
APPLICATION NUMBER: 08/461,271
FILING DATE:
APPLICATION NUMBER: 08/087,387
FILING DATE: 2-Jul-93
ATTORNEY/AGENT INFORMATION:
NAME: Stephen C. Macevicz
REGISTRATION NUMBER: 30,285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION
TELEPHONE: (415) 358-7855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (415) 358-7855
TELEPAX: (415) 358-7794
INFORMATION FOR SEO ID NO: 6: SEQUENCE CHARACTERISTICS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1143 CTTTTTTTTTT 1156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94306-2155
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGIH: 16 nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid
EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
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US-08-713-685A-6
US-08-713-685A-6
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US-08-689-856-6
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Gaps
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APPLICANT: Brown, Michael S.
APPLICANT: Briggs, Michael R.
APPLICANT: Briggs, Michael R.
APPLICANT: Wang, Xiaodong
APPLICANT: Wang, Xiaodong
APPLICANT: Goldstein, Joseph L.
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING
TITLE OF INVENTION: PROTEINS
NUMBER OF ENVENTION: PROTEINS
NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patenth Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/689,856
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/455,627
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12.4; DB 1;
Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                         NAME: NAKEMURA, Jackie N.
REGISTRATION NUMBER: 35,966
REGISTRATION NUMBER: 1XX-003/01 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-843-5000
TELEPHONE: 415-843-5000
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 nucleotides
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NUMBER OF SEQUENCES: 42
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/425,852
FILING DATE: 20-0CT-1989
PRIOR APPLICATION UNTRER: 33,330
FILING DATE: 30-MAR-1987
APPLICATION NUMBER: 33,081
FILING DATE: 30-MAR-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-373-124A-338/c
Sequence 338, Application US/08373124A
Patent No. 5646042
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Stinchcomb, Dan T. APPLICANT: Draper, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           471 GCAGGGGGGACT 484
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2 CTTTTTTTTTT 15
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Best Local Similarity
Lahes 13; Conserve
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Best Local Similarity
Matches 13; Conserva
                                                                            5256545-4/c
;Patent No. 5256545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO:33:
LENGTH: 16
                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO:4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5256545-33
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5256545-33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Serge: M. Gryaznov
TITLE OF INVENTION: Oligonucleotide clamps having diagnostic
TITLE OF INVENTION: and therapeutic applications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.9%; Score 12.4; DB 1; Length 16; Best Local Similarity 92.9%; Pred. No. 1.7e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                      0.9%; Score 12.4; DB 1; Length 16; 92.9%; Pred. No. 1.7e+02; rive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSE: Stephen C. Macevicz, Lynx Therapeutics
STRERT: Stephen Centre Drive
CITY: Foster City
STATE: California
CUNTRY: USA
ZIP: 94404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READBLE FORM:
MEDIUM TYPE: 5.25 inch diskette
COMPUTER: IBM compatible
COMPUTER: Windows 3.1/DOS 5.0
SOFTWARE: Microsoft Word for Windows, vers. 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/070,477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/713,685
FILING DATE:
APPLICATION NUMBER: 08/461,271
FILING DATE:
APPLICATION NUMBER: 08/087,387
FILING DATE: 2-041-93
ATORNEY/AGENT INFORMATION:
NAME: Stephen C. Macevicz
NAME: Stephen C. Macevicz
REGISTATION NUMBER: 30,285
                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 6, Application US/09070477 Patent No. 6048974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE DOCKET WINESE: 104
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 358-7855
TELEFAX: (415) 358-7794
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
TELEFAX: (512) 474-7577
INFORMATION FOR SEQ ID NO: 23
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                   471 GCAGGGGGAGT 484
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                                                                                                                                                                                                                                                                 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                    US-08-668-123-23
                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 274
US-09-070-477-6
                                                                                                                                                                                                                            Query Match
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Gaps
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APPLICANT: BROWN, MICHAEL S.;GOLDSTEIN, JOSEPH L.;RUSSELL,
;DAVID W.;SUDHOF, THOMAS C.;
;TITLE OF INVENTION: STEROL REGULATORY ELEMENTS
; VUMBER OF SEQUENCES: 42
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/425,852
; FILING DATE: 30-0000-1989
; PRIOR APPLICATION NUMBER: 33,330
; FILING DATE: 30-0000-1987
; FILING DATE: 30-0000-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 16;
                                                                                                                                                                                                                                                                                                                                                     Length 16;
APPLICANT: BROWN, MICHAEL S.; GOLDSTEIN, JOSEPH L.; RUSSELL, DAVID W.; SUDHOF, THOMAS C. TITLE OF INVENTION: STEROL REGULATORY ELEMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  METHODS AND COMPOSITIONS FOR
TREATMENT OF RESTENOSIS AND
CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                  0.9%; Score 12.4; DB 1;
92.9%; Pred. No. 1.7e+02;
cive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 12.4; DB 1;
Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: McSwiggen, James APPLICANT: Jarvis, Thale TITLE OF INVENTION: METHODS AN TITLE OF INVENTION: CANCER USINUMBER OF SEQUENCES: 2627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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1143 CTTTTTTTTTT 1156

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ADDRESSEE:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-173-124A-2049

US-08-173-124A-2049

Sequence 2049, Application US/08373124A

Patent No. 5646042

GENERAL INFORMATION:

APPLICANT: Stinchcomb, Dan T.

APPLICANT: Stinchcomb, James

APPLICANT: Stinchcomb, James

APPLICANT: Garvis, Thale

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR

TITLE OF SEQUENCES: 2627

CORRESPONDENCE ADDRESS:

ADDRESSEE: Lyon & Lyon

STORTE OF SEQUENCES: 2627

ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 8.0 rage
COMPUTER: IEM COMDALIBLE
COMPUTER: IEM COMDALIBLE
COPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PERFCE 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/373,124A
FILING DATE: JAHUARY 13, 1995
FRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/245,466
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/373,124A
FILING DATE: January 13, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/192,943
FILING DATE: FEbruary 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/986,422
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/986,422
FILING DATE: Abruguat 26, 1992
APPLICATION NUMBER: 07/987,132
FILING DATE: August 26, 1992
APPLICATION NUMBER: 209/035
FILING DATE: AUGUST 30, 209-160
TELECOMMUNICATION INFORMATION:
TELEPAX: 67-3510
INFORMATION FOR SEQ ID NO: 2047:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1118 GTTTAATTGAAAAA 1131
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4 GUUUUAUUGAAAA 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
US-08-373-124A-2047
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2047, Application US/08373124A
; Patent No. 5646042
; GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Stinchcomb, Dan T.
APPLICANT: McSwiggen, Venneth
APPLICANT: McSwiggen, James
APPLICANT: Graver General McSwiggen, James
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                               COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION INMEBR: US/08/373,124A
FILING DATE: January 13, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/373,124A
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: Rebraary 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/987,132
FILING DATE: ANGUST 26, 1992
ATORNEY AGENT INFORMATION:
MAME: WASDLIGY RICHARY
REFERENCE/DOCKET NUMBER: 209/035
TELEFONE: (213) 469-1600
TELEFONE: (213) 460-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                             SSSE: Lyon & Lyon
TT: 633 West Fifth Street
T: Suite 4700
Los Angeles
3: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 GGCAGGCAGTTGAG 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                        STATE: Californi
COUNTRY: U.S.A.
ZIP: 90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 278
US-08-373-124A-2047
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                                                                                                          STREET:
CITY: L
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Score 12.4; DB 1; Length 17; Pred. No. 1.9e+02; 0; Mismatches 1; Indels
0.9%; Score 12.4; DB 1; Length 17; 92.9%; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-435-628-338/C

Sequence 338, Application US/08435628

Patent No. 5817796

GENERAL INFORMATION:
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Draper, Kenneth
APPLICANT: McSwiggen, James
APPLICANT: Stinch S
                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,628
FILING DATE: 05-MAY-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/245,466
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/245,466
FILING DATE: February 7, 1994
APPLICATION NUMBER: 08/245,466
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 07/986,422
FILING DATE: AUGUST 07/936,422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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(213) 955-0440
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.9
Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS
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nucleic acid
                   Query Match
Best Local Similarity 92.9<sup>3</sup>
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single
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California
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COUNTRY: U.S.A.
ZIP: 90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH
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CITY: L
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| Sequence 30, Application US/08261822A
| Sequence 30, Application US/08261822A
| Sequence 30, Application US/0826182A
| Patent No. 5650553
| GENERAL INFORMATION:
| APPLICANT: Ecker, Joseph R. et al.
| APPLICANT: Ecker, Joseph R. et al.
| TITLE OF INVENTION: and Pathogens
| TITLE OF INVENTION: and Pathogens
| NUMBER OF SEQUENCES: 82
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Woodcock, Washburn, Kurtz, Mackiewicz & No. 5650553ris
| STREET: One Liberty Place, 46th floor
| CITY: Philadelphia | STREET: PATHOR | CORRESPONDENCE ADDRESSEES | CORRESPONDENCE 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .
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SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/261,822A
FILING DATE: 17-JUN-1994
CLASSIFICATION: 536
ATTONNEY/AGENT INFORMATION:
NAME: Beareful, Lori Y.
REGISTRATION NUMBER: 34,293
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEPHONE: (215) 568-3439
INFORMATION FOR SEQ ID NO: 30: SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                    FILING DATE: December 7, 1992
FILING DATE: December 7, 1996, 422
FILING DATE: August 26, 1992
ATTOCNEY AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 209/035
REFERENCE/DOCKET NUMBER: 209/035
TELEFONE: (213) 489-1600
TELEFAX: (213) 489-1600
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2049:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: Mucleic acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1118 GTTTAATTGAAAA 1131
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2 GUUUUAUUGAAAA 15
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Best Local Similarity 57.1:
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: linear
US-08-373-124A-2049
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
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-08-261-822A-30
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Gaps

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Sequence 2049, Application US/08435628 Patent No. 5817796 GENERAL INFORMATION:

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RESULT 282
US-08-435-628-2047

j Sequence 2047, Application US/08435628

patent No. 5817796

general INFORMATION:
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Draper, Kenneth
APPLICANT: Draper, Kenneth
APPLICANT: Dravis, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon

Lyon & Lyon

Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: California
COUNTRY: U.S.A.
ZIE: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 8torage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURKENT AFFLICATION NUMBER: US/08/435,628
FILING DATE: 05-MAY-1995
CLASSIFICATION: 514
FRICA APPLICATION DATA:
APPLICATION NUMBER: 08/373,124
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/936,422
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
FILING DATE: August 26, 1992
APTCORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               209/035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 209/
TELECOMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: 67-3510
INFORMATION FOR SEQ ID NO: 2047:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                     E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       32,327
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REGISTRATION NUMBER: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                       Los Angeles
: California
XY: U.S.A.
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APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Draper, Kenneth
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENCES AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & IT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: Ward Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,628
FILING DATE: 05-MAY.1995
CLASSIFTCATION DATA:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/373,124
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/245,466
FILING DATE: February 7, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/987,132
FILING DATE: August 26, 1992
APPLICATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,927
FELENCOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 284
US-08-485-611A-9
Sequence 9, Application US/08485611A
Setence 9, Application US/08485611A
RENERAL INFORMATION:
APPLICANT: MAYEL, Jr., Rich B.
APPLICANT: Gamper, Howard B.
APPLICANT: Kutyavin, Igor V.
                                                                                                                                                                                                                                                                                     iB: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1118 GTTTAATTGAAAA 1131
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GUUUUAUUGAAAAA 15
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SS: single
linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 8; Conserva
                                                                                                                                                                                                                                                                                                                                                        CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
TOPOLOGY: ]
US-08-435-628-2049
                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U
                                                                                                                                                                                                                                                                                                           STREET:
STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              g
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1118 GTTTAATTGAAAA 1131 |::: |::||||||| 4 GUUUUAUUGAAAA 17

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RESULT 283 US-08-435-628-2049

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0.9%; Score 12.4; DB 1; Length 17; 50.0%; Pred. No. 1.9e+02; tive 6; Mismatches 1; Indels
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US-08-962-162-119
| Sequence 119, Application US/08985162
| Sequence 119, Application US/08985162
| GENERAL INFORMATION:
| APPLICANT: Akhtar, Saghir
| APPLICANT: McSwiggen, James
| TILLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT ITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT ITLE OF INVENTION: TO LEVELS OF EPIDERMAL GROWTH
| TITLE OF INVENTION: TO LEVELS OF EPIDERMAL GROWTH
| TITLE OF INVENTION: FACTOR RECEPTORS
| NUMBER OF SEQUENCES: 1877
| CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                             APPLICANT: ACCOUNTING SAGAIL
APPLICANT: ACADIAN SAGAIL
APPLICANT: Fell, Patricia
APPLICANT: Fell, Patricia
APPLICANT: Fell, Patricia
APPLICANT: McSwiggen, James
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT
TITLE OF INVENTION: TO LEVELS OF EPIDERMAL GROWTH
TITLE OF INVENTION: FACTOR RECEPTORS
NUMBER OF SEQUENCES: 1877
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon
STREET: 633 Weet Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: CALLLOLLING
COUNTRY: CALLLOLLING
ZIP: 90071-206
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FRASTSEQ for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/985,162
FILING DATE: 04 December 1997
CLASSIFTCATION STATE
APPLICATION NUMBER: 00/036,476
FILING DATE: 31 January 1997
ATTORNEY ACENT THORMATION:
NAME: WALDLY RICHARD A
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
TELECOMMUNICATION INFORMATION:
TELECHNUM: (213) 9499-1600
                                                                                                                                                ; Sequence 118, Application US/08985162
; Patent No. 6057156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1099 CGTAATTATGTAGT 1112
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Best Local Similarity 50.05
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
US-08-985-162-118
                                                                                                                      US-08-985-162-118
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81.2%; Pred. No. 1.9e+02;
iive 1; Mismatches 2; Indels
APPLICANT: Gall, Alexander A.
APPLICANT: Petrie, Charles R.
APPLICANT: Tabone, John C.
APPLICANT: Tabone, John C.
TITLE OF INVENTION: CROSSLINKING OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: Klein & Szekeres
STREET: Al99 Campus Drive, Suite 700
CITY: Irvine
STREE: CA
                                                                                                                                                                                                                                                                                                                                                                                                    CUUNTY: USA
ZIPE: 92715
COUNTY: USA
ZIPE: 92715
COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,611A
FILING DATE: 07-JUN-1995
CLASSIFICATION: 514
PRICK APPLICATION DATA:
APPLICATION NUMBER: US 08/226,949
FILING DATE: 27-JUN-1994
RAIDA APPLICATION NUMBER: US 08/011,482
FILING DATE: CS-JNN-1994
RAIDA APPLICATION DATA:
APPLICATION NUMBER: US 08/34,490
FILING DATE: US 08/34,490
FILING DATE: US 08/34,490
FILING APPLICATION DATA:
APPLICATION NUMBER: US 08/34,490
FILING APPLICATION DATA:
APPLICATION NUMBER: US 07/353,857
FILING DATE: 21-AUG-1991
ATTORNEY/AGENT INFORMATION:
NUMBE: SZEGAFES, GDAD: L.
N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Szekeres, Gabor L. REGISTRATION NUMBER: 28,675 REFERENCE/DOCKET NUMBER: 491 TELECOMMUNICATION: 714-854-5502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1147 TTTTCTTTTGGAAGT 1162
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INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 TTTTCTTTTYGGGGGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.9%
Best Local Similarity 81.2%
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-485-611A-9
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                                                                                                                                                                                                                                                                                            Sequence 79, Application US/09017974

Sequence 79, Application US/09017974

Patent No. 6288042

GENERAL INFORMATION

APPLICANT: Rando, Robert F.

APPLICANT: Hogan, Michael B.

APPLICANT: Hogan, Michael B.

APPLICANT: Gosum, Paul A.

TITLE OF INVENTION: Anti-Viral Guanosine-Rich

TITLE OF INVENTION: Tetrad Forming Oligonucleotides

NUMBER OF SQUENCES. 88

CORRESPONDENCE ADDRESS:

ADDRESSE: Conley, Rose & Tayon, P.C.

STREET: Houston

CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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illarity 92.9%; Pred. No. 1.9e+02;
Conservative 0; Mismatches 1;
                                                                                Best Local Similarity 71.4%; Fred. No. 1.9e+02; Matches 10; Conservative 3; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: TEXAS

STATE: TEXAS

CONDURKY: U.S.A.

ZIP: 77002-232

COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS Word 97 (saved as .txt file)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/017,974
FILING DATE: 04-FEB-97
APPLICATION NUMBER: 60/037,374
FILING DATE: 04-FEB-97
APPLICATION NUMBER: 60-DC-97
ATTORNEY/AGENT INFORMATION:
NUMBER: 09-DEC-97
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-682-255A-79

Sequence 79, Application US/08682255A

Patent No. 6323185

GENERAL INFORMATION:

APPLICANT: Rando, Robert F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 33,962
REGISTRATION NUMBER: 31,962
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/239-8010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA (genomic) US-09-017-974-79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              762 GTGGCGGGTGGATG 775
                                                                                                                                               56 CTCCTCAATTACCC 69
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4 CUCCUCAAUGACCC 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 713/238-8010
TELEPAX: 713/238-8008
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                              RESULT 288
US-09-017-974-79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 289
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; Sequence 95, Application US/0899809A
; Patent No. 610380
; GERERAL INFORMATION:
APPLICANT: GARVIS, THALE
; APPLICANT: MCSWIGGEN, JAMES A.
APPLICANT: STINCHCOMB, DAN T.
TITLE OF INVENTION: ENCYMATIC NUCLEIC ACID TREATMENT OF DISEASES
; TITLE OF INVENTION: OR CONDITIONS REHATED TO LEVELS OF C-FOS
; TITLE OF INVENTION: OR CONDITIONS
FILE REPERENCE: 231/175
; CURRENT APPLICATION NUMBER: US/08/998,099A
; CURRENT FILING DATE: 1997-11-2-4
; EARLIER APPLICATION NUMBER: 60/037,658
; EARLIER APPLICATION NUMBER: 08/373,124
; EARLIER PILING DATE: 1995-01-13
; EARLIER PILING DATE: 1995-01-13
; EARLIER PILING DATE: 1995-01-13
; SARLIER PILING DATE: 1995-01-13
; SARLIER PILING DATE: 1995-01-13
; SARLIER PILING DATE: 1995-01-13
; SOFTWARE: FASEQ ID NOS: 375
; SOFTWARE: FASESQ for Windows Version 3.0
; SEQ ID NO 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%; Score 12.4; DB 1; Length 17; 50.0%; Pred. No. 1.9e+02; tive 6; Mismatches 1; Indels
                                                                                                                                                                                                     COUNTY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OOPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/985,162
FILING DATE: 04 December 1997
CLASSIFICATION: 514
PRIGR APPLICATION DATA:
APPLICATION NUMBER: 60/036,476
FILING DATE: 31 January 1997
ATTORNEY APPLICATION NUMBER: 83,327
REFERENCE/DOCKT NUMBER: 23,327
REFERENCE/DOCKT NUMBER: 23,327
TELECHONE: (213) 489-1600
TELECHONE: (213) 955-0440
LYON & Lyon to Lyon cet 2BET: Suite 4700
': Los Angeles
': California
': U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1099 CGTAATTATGTAGT 1112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match: 0.9%
Best Local Similarity 50.0%
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nuclear
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-985-162-119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 287
US-08-998-099-95
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                                                                                                                                               CITY: LOS
STATE: Ca
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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0.9%; Score 12.4; DB 1; Length 17; 71.4%; Pred. No. 1.9e+02;
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Fatent No. 634638
GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: Bcsobedo, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Scobedo, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                   ADDRESSE: Lyon & Lyon STREET: 633 West Fifth Street STREET: 633 West Fifth Street STREET: Los Angeles CTIY: Los Angeles CALIFORNIA COUNTRY: U.S.A. ZIP: 90071-2066 COUNTRY READABLE FORM: MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION DATA:
APPLICATION NUMBER: 0/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELEFHONE: (213) 955-0440
                                                                                              GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E. Lyon & Lyon
633 West Fifth Street
Suite 4700
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTER.STICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 UUGGAACUAAAGCA 14
                                                                                                                               8502
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Best Local Similarity 71.49
Matches 10; Conservative
      TITLE OF INVENTION: TREA TITLE OF INVENTION: COND TITLE OF INVENTION: OF V TITLE OF INVENTION: GROWN THERESPONDENCE ADDRESS:
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STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: ]
US-08-584-040-2256
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US-08-584-040-2547
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STREET: 63
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0.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 1.9e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2256, Application US/08584040
Patent No. 6346398
GENERAL INFORMATION:
APPLICANT: Pavoco, Pamela
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Becobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
                                                                                                                                                                                                                                                                                                                                                                              CITEL: 1 COUTINGS, SUITE 100 CITES: 1 COUNTY: 
                                                                                                                                  APPLICANT: Pommier, Byves
APPLICANT: Maxumder, Abhijit
TITLE OF INVENTION: Anti-Viral Guanosine-Rich
TITLE OF INVENTION: Oligonucleotides
CORRESPONDENCE: 87
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Conley, Rose & Tayon, P.C. STREET: 600 Travis, Suite 1850 CITY: Houston
      Fennewald, Susan
Zendegui, Joseph G.
Ojwang, Joshua O.
Hogan, Michael E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/238-8010
TELEFAX: 713/238-8018
INFORMATION FOR SEQ ID NO: 79:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: DNA (genomic)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-584-040-2256
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:

APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: Bachedo, James
APPLICANT: Eschedo, Jame TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: GROWTH FACTOR
MUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: State 4700
CHIV: LOS ANGELSS
CHIVE AN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
                                                                                                ATIONALY ASELL TREATED.

REGISTRATION NUMBER: 32,327

REGISTRATION NUMBER: 32,327

REFERENCE/DOCKET NUMBER: 32,327

TELECOMMUNICATION INFORMATION:

TELEPHONE: (213) 489-1600

TELEFAX: (213) 955-0440

SEQUENCE CHARACTERISTICS:

LENGTH: 17 base pairs

TYPE: nucleic acid

STARMEDNESS: single

STARMEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-584-040-2549
; Sequence 2549, Application US/08584040
; Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       218/064
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APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 60/005,974
FILING DATE: 0ctobez 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2549:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (213) 489-1600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 CUUUUUUUUUUUU 16
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COUNTRY: U.S.A.
ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dp
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US-08-584-040-2548
IS-08-584-040-2548
Sequence 2546 Application US/08584040
SEQUENCE Stindscomb, Dan T.
APPLICANT: Bacobedo, James
APPLICANT: Bacobedo, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWIH FACTOR
TITLE OF INVENTION: GROWIH FACTOR
TITLE OF INVENTION: GROWIH FACTOR
NUMBER OF SEQUENCES: B502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STREET: California
STREET: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 7.1%; Pred. No. 1.9e+02; 1; Conservative 12; Mismatches
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ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
                                                                                                              MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFWWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
APTORNEY, AGENT INFORMATION:
NAME: WAPDLICATION:
NAME: WAPDLICATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECHONE: (213) 955-0440
TELEPHONE: (213) 955-0440
TELERAX: 67-3510
INFORMATION FOR SEQ ID NO: 2547:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
INFORMATION FOR SEQ ID NO: 2547:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TERES: ALTORATE AND ASSES SINGLE
                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1143 CTTTTTTTTTTT 1156
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STRANDEDNESS: Sir
mobology: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 1; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-584-040-2547
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17 base pairs

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(213) 489-1600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELER: (213) 955-044C
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15 AAACCAAACCCTGC 2
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Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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                                                                  US-08-584-040-6008/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-584-040-6008
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                                           RESULT 295
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                                                                                                     Query Match 0.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 7.1%; Pred. No. 1.9e+02; Matches 1; Conservative 12; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Parco, Pamela
APPLICANT: Parco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T
APPLICANT: Stinchcomb, Dan T
APPLICANT: Escobedo, Jame
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: LYON & LYON
STREET: 633 West Fifth Street
STREET: STATE 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITYEL: LOS ATGGELES
CITY: LOS ATGGELES
COUNTRY: U.S.A.
ZIP: 20071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
COMPUTER: IBM COMPATIBLE FO. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION MUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                Sequence 2550, Application US/08584040 Patent No. 6346398 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        218/064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 218
PELECOMONIACATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                               1143 CTTTTTTTTTTT 1156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                        ; TYPE: nucleic acid
; STRANDEDRESS: single
; TOPOLOGY:
US-08-584-040-2549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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US-08-584-040-2550
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1143 CTTTTTTTTTTT 1156

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Gaps
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Sequence 600, Application US/08584040

Patent No. 6346398

GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: Bacodo, Jaime
APPLICANT: Escobed, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWITH PACTOR
TITLE OF INVENTION: GROWITH FACTOR
TITLE OF INVENTION: GROWITH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon

ADDRESSEE: Lyon & Lyon

ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 12.4; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READMBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REFIRENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION: NUMBER: 218/064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-584-040-6009/c; Sequence 6009, Application US/08584040; Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 218/064
                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Houston
STATE: Texas
COUNTRY: U.S.A.
ZIP: 7702-2912
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTER: MS Windows 95
SOFTWARE: MS Word 97 (saved as .txt file)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Anti-Viral Guanosine-Rich Oligonucleotides NUMBER OF SEQUENCES: 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Conley, Rose & Tayon, P.C.
STREET: 600 Travis, Suite 1850
                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: U1/19/12,1996
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/001,135
FILING DATE: U1/13,1995
APPLICATION NUMBER: 60/001,26
FILING DATE: SEPTEMBER: 1994
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHANE: (213) 955-0440
                                                                                                                                                  ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fennewald, Susan
Zendegui, Joseph G.
Ojwang, Joshua O.
Hogan, Michael E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 79, Application US/09429130 Patent No. 6355785 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pommier, Eyves
Mazumder, Abhijit
60/015,714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Rando, Robert F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 884 TCCAGGAGCTGCGG 897
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
STREET: Sulce
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 298
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Zwick, Michael G.
APPLICANT: Zwick, Michael G.
APPLICANT: Edington, Brent B.
APPLICANT: McSwiggen, James A.
APPLICANT: McSwiggen, James A.
APPLICANT: Guo, Lining
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Young, Scott A.
APPLICANT: Folkerts, Otto
APPLICANT: Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR
TITLE OF INVENTION: MODULATION OF GENE EXPRESSION
TITLE OF INVENTION: IN PLANTS
NUMBER OF SEQUENCES: 1263
              METHOD AND REAGENT FOR THE TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                                               CITY: Los Angeles
STATE: California
COUNTRY: U.S.A..
ZIP: 30071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: OBLUAY 11, 1996
CLASSIFICATION 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60,005,974
FILING DATE: October 26, 1995
ATTORNEY AGENT INFORMATION:
NAME: WARDING, RIGHARD 4.
REGISTRATION NUMBER: 32,337
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMINICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 139, Application US/08679645
Patent No. 6350934
GENERAL INFORMATION:
                                                                                                                                                  GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                             3: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
          TITLE OF INVENTION: METHOD TITLE OF INVENTION: CONDITION: TITLE OF INVENTION: OF VASCITATE OF INVENTION: GROWTH OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 67.3510
INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23 AAACCAAACCCAGC 36
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Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyo
                                                                                                                                                                                                                                                                                                                             Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
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US-08-679-645-139/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
US-08-584-040-6009
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                                                                                                                                                                                                                                                                                      STREET:
                                                                                                                                                                                                                                                                                                                             STREET:
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APPLICANT: Karpeisky, Alex
APPLICANT: Adamic, Jasenka
APPLICANT: Sweedler, David
APPLICANT: Zinnen, Sharinen, Sharin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        us-uy-suv-stant to the factor of the factor 
                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 1.9e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                   Score 12.4; DB 1; Length 17;
Pred. No. 1.9e+02;
); Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 1999-12-19
PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR PLILING DATE: 1997-11-05
PRIOR PLILING DATE: 1991-04-29
PRIOR FILING DATE: 1998-04-29
PRIOR FILING DATE: 1998-11-04
PRIOR PLILING DATE: 1998-11-04
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR APPLICATION NUMBER: US 09/181,511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 409, Application US/09474432B
Patent No. 6528640
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1144 TITITITICITITG 1157
                                                                                                                                        llarity 92.9%;
Conservative
                                                                                                                                                                                                                                                                                      1144 TITTTTTTTTTG 1157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Beigelman, Leo
Burgin, Alex
Beaudry, Amber
                                                                                            Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 301
US-09-474-432B-409
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APPLICANT:
APPLICANT:
US-09-788-338-3
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Sequence 3, Application US/09788338

Patent No. 648596

GENERAL INFORMATION

APPLICANT: MURAMATSU, TAKAMICHI

APPLICANT: KIYAMA, MASAHARU

APPLICANT: KIYAMA, MASAHARU

APPLICANT: KIYAMA, MASAHARU

TITLE OF INVENTION: NUCLEIC ACID SAMPLE FOR RARE

TITLE OF INVENTION: BYPRESSED GENES AND ANALYZING METHOD USING THE PREPARED

TITLE OF INVENTION: NUCLEIC ACID SAMPLES THEREBY

TITLE OF INVENTION: NUCLEIC ACID SAMPLES THEREBY

TITLE OF INVENTION: NUCLEIC ACID SAMPLES THEREBY

TITLE OF LINGBRICATION NUMBER: US/09/788,338

CURRENT FILING DATE: 1999-05-18

PRIOR FILING DATE: 1999-05-18

PRIOR FILING DATE: 1999-05-20

NUMBER OF SEQ ID NOS: 4

SOFTWARE: PATENTIN VOE: 2.0
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FRATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR AFPLICATION INDRERS: 08/682,255
FILING DATE: -CUKNOWNS
APPLICATION NUMBER: 60/001,505
FILING DATE: 19-UULY-95
APPLICATION NUMBER: 60/014,007
FILING DATE: 25-MARCH-96
APPLICATION NUMBER: 60/014,688
FILING DATE: 19-WARCH-96
APPLICATION NUMBER: 60/016,271
FILING DATE: 17-APRIL-96
ATTORNEY/AGENT INFORMATION:
NAME: MODENIAL, C. SEEVEN
REFERENCE/DOCKET NUMBER: 33,962
REFERENCE/DOCKET NUMBER: 33,962
REFERENCE/DOCKET NUMBER: 31,962
REFERENCE/DOCKET NUMBER: 1472-06214
TELLEPHONE: 713/238-8010
TELLEPHONE: 713/238-8018
INFORMATION FOR SECILO NO: 79:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                APPLICATION NUMBER: US/09/429,130
FILING DATE: 28-Oct-1999
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
SEQUENCE DESCRIPTION: SEQ ID NO: 79:
                                                                                                                                                                                                                                          25-MARCH-96
19-MARCH-96
                                                                                                                                                                                                                                                                                                                                       17-APRIL-96
23-APRIL-96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
         CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          762 GIGGCGGGTGGATG 775
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Best Local Similarity
Matches 13; Conservi
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US-09-788-338-3
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LENGTH: 17
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TITLE OF INVENTION: Mucleotide triphosphate and their incorporation into oligomucle;
CURRENT APPLICATION WHOBER: US/09/474,432B

CURRENT FILING DATE: 1999-12-19

PRIOR APPLICATION NUMBER: US 60/064,866

PRIOR FILING DATE: 1999-12-10

PRIOR FILING DATE: 1998-04-29

PRIOR FILING DATE: 1998-04-29

PRIOR FILING DATE: 1998-11-04

PRIOR FILING DATE: 1998-11-04

PRIOR FILING DATE: 1999-04-28

PRIOR FILING DATE: 1999-04-28

SEQ ID NOS: 1526

SOFTWARE: Patentin version 3.0

LENGTH: 17

TYPE: RA

ORGANISM: Homo sapiens

US-09-474-432B-557
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.9%; Score 12.4; DB 1; Length 17; 21.4%; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.9%; Score 12.4; un
Best Local Similarity 21.4%; Pred. No. 1.96+
Matches 3; Conservative 10; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 815, Application US/09474432B
Patent No. 6528640
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1112 TTTTCTGTTTAATT 1125
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Best Local Similarity 92.99
Matches 13; Conservative
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US-09-474-432B-815
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US-09-371-772B-801
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GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Beigelman, Leo
APPLICANT: Beigelman, Leo
APPLICANT: Beigelman, Leo
APPLICANT: Beaudry, Amber
APPLICANT: Beaudry, Amber
APPLICANT: Sameni, Jasenka
APPLICANT: Sameni, Shawn
FILE REFERENCE: MBHB00-831-B (247/276)
CURRENT APPLICATION NUMBER: US 60/064, B66
FRIOR APPLICATION NUMBER: US 60/064, B66
FRIOR APPLICATION NUMBER: US 60/084, 727
PRIOR APPLICATION NUMBER: US 09/186, 675
FRIOR APPLICATION NUMBER: US 09/186, 675
FRIOR FILING DATE: 1998-04-29
FRIOR FILING DATE: 1998-04-29
FRIOR FILING DATE: 1999-04-28
FRIOR FILING DATE: 1999-04-28
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                                                                                                                                                                                                                                                  0.9%; Score 12.4; DB 1; Length 17; 85.7%; Pred. No. 1.9e+02; tive 1; Mismatches 1; Indels
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APPLICANT: Beigelman, Leo
APPLICANT: Burgin, Alex
APPLICANT: Beaudry, Amber
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; Sequence 421, Application US/09474432B
; Patent No. 6528640
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        PRIOR FILING DATE: 1999-04-28
NUMBER OF SEQ ID NOS: 1526
SOFFWARE: PatentIn version 3.0
SEQ ID NO 409
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1 GGAGGAAUGCCGAG 14
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Adamic, Jasenka
Sweedler, David
Zinnen, Shawn
                                                                                                                                                                                                                                            Query Match
Best Local Similarity 85.7
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
CORGANISM: Homo sapiens
US-09-474-432B-421
                                                                                                                                              TYPE: RNA
ORGANISM: Homo sapiens
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
                                                                                                                  LENGTH: 17
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GENERAL INFORMATION:

APPLICANT: BLOEZWHE Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: Patrochoch, Dan
APPLICANT: Bscobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00, 876-J (237/198)
CURRENT APPLICATION NUMBER: US 60/005, 974
FRICK APPLICATION NUMBER: US 60/005, 974
FRICK APPLICATION NUMBER: US 68/584, 040
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: ParentIn version 3.0
SEQ ID NO 1072
LENGTH: 17
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APPLICANT: StinchComb, Dan
APPLICANTON: Escobado, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
FILE REFERENCE: MBHB00,876-J (237/198)
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT PAPLICATION NUMBER: US /09/371,772B
FRIOR FILING DATE: 1999-08-10
PRIOR FILING DATE: 1999-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Parentin version 3.0
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0.94; Source ...,

Best Local Similarity 7.18; Pred. No. 1.96+02;

1. Conservative 12; Mismatches 1;
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Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Pavco, Pam
APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 1074, Application US/09371772B ; Patent No. 6566127
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US-09-371-772B-1073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Homo sapiens US-09-371-772B-1072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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US-09-371-772B-1074
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                                                                                              APPLICANT: Pavco, Pam
APPLICANT: Barco, Pam
APPLICANT: Barco, Jim
APPLICANT: Stinchomb, Jan
APPLICANT: Stinchomb, Jan
APPLICANT: Stinchomb, Jan
APPLICANT: Stinchomb, Jan
APPLICANT: Barcobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHBOO, 376-JU (237/198)
CURRENT APPLICATION NUMBER: US 60/005, 974
PRIOR APPLICATION NUMBER: US 60/005, 974
PRIOR APPLICATION NUMBER: US 08/584, 040
PRIOR FILING DATE: 1996-10-26
PRIOR FILING DATE: 1996-10-8
SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 801
LIBRICH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Becobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
CURRENT APPLICATION WMBER: US/09/371,772B
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEQ ID NO 1071
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match

0.9%; Score 12.4; DB 1; Length 1
Best Local Similarity 71.4%; Pred. No. 1.9e+02;
Matches 10; Conservative 3; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
                                                                                   APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1071, Application US/09371772B
Patent No. 6566127
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Sequence 801, Application US/09371772B
Patent No. 6566127
GENERAL INFORMATION:
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Best Local Similarity 7.1%;
Matches 1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CRGANISM: Homo sapiens
US-09-371-772B-1071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 307
US-09-371-772B-1072
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US-09-371-772B-1071
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APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: MENBER: US/09/371,728
CURRENT APPLICATION NUMBER: US/09/371,728
FRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEQ ID NO 2946
LENGTH::
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE REFERENCE: MBHB00, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 68/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
NUMBER OF SEQ ID NOS: 144225
SOFTWARE: Patent In version 3.0
SEQ ID NO 5053
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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Pred. No. 1.9e+02;
1; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 5053, Application US/09371772B
Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
Ribozyme Pharmaceuticals, Inc.
Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1
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ilarity 85.7%;
Conservative 1
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Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: Mus sp. US-09-371-772B-2846
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ÚS-09-371-772B-5053
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US-09-371-772B-5053
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US-09-371-772B-2845/c

j Gequence 2445, Application US/09371772B

gequence 2445, Application US/09371772B

gequence 2445, Application US/09371772B

general No. 6566127

GENERAL INFORMATION:

APPLICANT: Pavco, Pam

APPLICANT: Pavco, Pam

APPLICANT: Stinchcomb, Dam

APPLICANT: Stinchcomb, Dam

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION: Wethod and Reagent for the Treatment of Diseases

CURRENT PILING DATE: 199-10-10-26

PRIOR PELICATION NUMBER: US 60/005,974

PRIOR PELICATION NUMBER: US 60/005,974

PRIOR APPLICATION NUMBER: US 60/005,974

PRIOR APPLICATION NUMBER: US 68/584,040

NUMBER OF SEQ ID NOS: 14225

SOFTWARE: PatentIn version 3.0

SEQ ID NO 2845

LENGTH: 17
                                                                                                  APPLICANT: Stinchcomb, Dan
APPLICANT: Sacobedo, Jaine
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REPERENCE: MBHB00, 876-J (237/198)
CURRENT APPLICATION NUMBER: US 60/005, 974
PRIOR APPLICATION NUMBER: US 60/005, 974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-001-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1074
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.9%; Score 12.4; DB 1; Length 17; 7.1%; Pred. No. 1.9e+02; tive 12; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
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                                    APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 311
US-09-371-772B-2846/c
Sequence 2846, Application US/09371772B
; Patent No. 6566127
; GENERAL INFORMATION:
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Best Local Similarity 92.9
Matches 13; Conservative
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                                                                                         McSwiggen, Jim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 1; Conserv
              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ), ORGANISM: Mus sp. US-09-371-772B-2845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-371-772B-1074
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APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stochedo, Jaime
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE REPERENCE: MHBB00, 876-J (237/198)
CURRENT PILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
RICHARRE: PAPELICATION NUMBER: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 6554
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GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Plant Genes for Sensitivity to Ethylene
TITLE OF INVENTION: and Pathogens
NUMBER OF SEQUENCE: 82
CORRESPONDENCE ADDRESS:
ADDRESSE: Woodcock, Washburn, Kurtz, Mackiewicz & Norris
STREET: One Liberty Place, 46th floor
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.9%; Score 12.4; DB 1; Length 17; 78.6%; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTR: ...
ZIF: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Extent Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/07744A
TTTING DATE: 15-UUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/261,822
FILING DATE: June 17, 1994
ATTORNEY/ACENT INFORMATION:
NAME: Beardell, Lori Y.
REGISCHARATION NUMBER: 34,293
TELECOMMUNICATION INFORMATION:
TELEDHONE: (215) 568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (215) 568-31C
TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               550 CTGGCAGGCATGCA 563
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Best Local Similarity 78.69
Matches 11; Conservative
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TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Homo sapiens
US-09-371-772B-6554
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCT-US95-07744A-30/C
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GENERAL INCORMATION:
GENERAL INCORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Racwigaen, Jim
APPLICANT: Racwigaen, Jim
APPLICANT: McSwigaen, Jim
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00, 876-J (237/198)
FILE REFERENCE: MUMBER: US 60/005,974
PRIOR FILING DATE: 1995-00-10
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 5055
LENGTH: 17
                                        APPLICANT: MCSW-199En, Jim
APPLICANT: StinchComb, Dan
APPLICANT: StinchComb, Jan
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
CURRENT APPLICATION NUMBER: US/09/371,72B
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-01-03
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 12.4; DB 1; Length 17; Pred. No. 1.9e+02; 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-371-772B-6554
; Sequence 6554, Application US/09371772B
; Patent No. 6566127
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5055, Application US/09371772B Patent No. 6566127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1313 AGCCAGGTGCTTTT 1326
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  687 IGGGAGCCAGCGGC 700
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Best Local Similarity 85.73
Matches 12, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-371-772B-5055
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US-09-371-772B-5055
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                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 5054
                           APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Draper, Kenneth G.
APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Gustofson, John
APPLICANT: Gustofson, John
APPLICANT: Stinchcomb, Dan T.
TITLE OF INVENTION: OF APTHRITIC CONDITIONS
NUMBER OF SEQUENCES: 1151
CORRESPONDENCE ADDRESS:
ADDRESSER: Lyon
STREET: Suite 4700
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/354,920
FILING DATE: December 13, 1994
APPLICATION NUMBER: 08/152,487
FILING DATE: No. 5612215ember 12, 1993
APPLICATION NUMBER: 07/989,848
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FaetSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,850
FILING DATE: February 17, 1995
0372/09696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 589, Application US/08390850 Patent No. 5612215 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 589:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  441 AAAGTTGCTGAAGTTTG 457
                    TELECOMMUNICATION INFORMATION:
TELEPHONE: 212/527-7700
TELEFAX: 212/753-6237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17 AAATTTGATGAAGTATG 1
     REFERENCE/DOCKET NUMBER:
                                                                      TELEFAX: 212/753-6237
TELEX: 236687
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 633 West Fi
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                               IMMEDIATE SOURCE:
CLONE: 711+1GTN
                                                                                                                                                                                                                                               TOPOLOGY: line MOLECULE TYPE: c ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JS-08-390-850-589/c
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                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: MISSING FIRAL MACRELICATE, AND APPLICANT: MISSING FIRAL MACRELICANT: MCSWiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stindshowb, Dan
APPLICANT: Stindshowb, Dan
APPLICANT: Stochedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Merels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHBOO, 976-0 (237/197)
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR PAPLICATION NUMBER: US 60/005,974
PRIOR PAPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEQ ID NO 5055
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                                                      Gaps
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Sequence 49, Application US/08281940

Fatent No. 5589330

GENERAL INFORMATION:

TITLE OF INVENTION:

TITLE OF INVENTION:

NUMBER OF SEQUENCES: 65

CORRESPONDENCE ADDRESS:

ADDRESSEE: DARBY & DARBY P.C.

STREET: 805 THIRD AVENUE

CITY: NEW YORK

STRATE: NEW YORK

STRATE: NEW YORK
Query Match 0.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 1.9e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 2.1e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/281,940
                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Ribozyme_Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                  US-09-371-772B-5055/c
; Sequence 5055, Application US/09371772B
; Patent No. 6566127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1027 CAAAGTGCAGCTGACTC 1043
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ATTORNEY/AGENT INFORMATION:
NAME: LUDWIG, S. PETER
REGISTRATION NUMBER: 25351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17 CÁAAAAGCAGCTGGCTC 1
                                                                                                  648 CCCCCAAGACCTGG 661
                                                                                                                                                 17 ccaccaagaccres 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ÚS-09-371-772B-5055
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Length 17;
                                                                                                                                           Sequence 59.2, Application US/08390850
; Sequence 59.2, Application US/08390850
; Patent No. 5612215
; GENERAL INFORMATION:
APPLICANT: Draper, Kenneth G.
APPLICANT: Pavco, Pamela
APPLICANT: Gustofson, John
APPLICANT: Stinchcomb, Dan T.
TITLE OF INVENTION: OF ARTHRITIC CONDITIONS
ITTLE OF INVENTION: OF ARTHRITIC CONDITIONS
NUMBER OF SEQUENCES: 1151
CORRESPONDENCE ADDRESS:
ADDRESSE: ADDRESSE:
ADDRESSE: ADDRESSE:
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.9%; Score 12.2; DB 1;
Best Local Similarity 58.8%; Pred. No. 2.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MADLIN TYPE: SCORAGGE
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,850
FILING DATE: February 17, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/354,920
FILING DATE: December 13, 1994
APPLICATION NUMBER: 08/152,487
FILING DATE: No. 5612215ember 12, 1993
APPLICATION NUMBER: 07/989,848
FILING DATE: No. 5612215ember 12, 1993
APPLICATION NUMBER: 07/989,848
FILING DATE: No. 5612215ember 12, 1993
APPLICATION NUMBER: 32,327
ATTORNEY/AGENT INFORMATION:
NAME: WALCATION INFORMATION:
TELEBRANCE/DOCKET NUMBER: 21,1084
TELEBRANCE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 322
US-08-435-634-589/c
'Sequence 589, Application US/08435634
'Patent No. 5731295
'GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1056 CCCTGGCCTTCCCATCA 1072
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364 TICTIGGGGCCCAGCT 380
                                                17 Trcrreservaccascr 1
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: Sulte .
CITY: Los Angeles
cuarte: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90071
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Ca
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                                                                                                                                                                    Query Match 0.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 2.1e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: McSwiggen, James
APPLICANT: Gustofson, John
APPLICANT: Glustofson, John
TITLE OF INVENTION: METHOD AND REAGENT FOR TREATMENT
TITLE OF INVENTION: OF ARTHRITIC CONDITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 5.5" Diskette, 1.44 Mb
MEDIUM TYPE: storagel
COMPUTER: IBM COMPALIAN
COMPUTER: IBM COMPALIAN
COMPUTER: IBM COMPALIAN
COMPUTER: BESELEG VERSION 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,850
FILING DATE: PEDEUARY 17, 1995
PRIOR APPLICATION NUMBER: 08/354,920
FILING DATE: December 13, 1994
APPLICATION NUMBER: 08/152,487
FILING DATE: No. 5612215ember 12, 1993
APPLICATION NUMBER: 07/989,848
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: DECEMBER 7, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 590, Application US/08390850
Patent No. 561215
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 1151
CORRESPONDENCE ADDRESS:
ADDRESSEE: 1yon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRX: U.S.A.
ZIP: 90071.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: WALDING, EACHARD
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 311/0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
TELEK: 67-3510
INFORMATION FOR SEQ ID NO: 590:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDENDESS: single
                                                                                                                                                                                                                                                                            365 TCTTGGGGGCCCCAGCTT 381
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Draper, Kenneth G.
Pavco, Pamela
                                                                                                                                                                                                                                                                                                                       17 refréégraaccagerr 1
     ; SEQUENCE CHARACTERISTICS:
; LENGTH: 17 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-390-850-589
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US-08-390-850-590/c
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APPLICANT:
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| Sequence 592, Application US/08435634
| Sequence 592, Application US/08435634
| Patent No. 5731295
| GENERAL INPORMATION:
| APPLICANT: Draper, Kenneth G. APPLICANT: Draper, Memela APPLICANT: McSwiggen, James APPLICANT: Guencheson, James APPLICANT: McSwiggen, James McGuntant McSwiggen, James McGuntant McSwiggen, McSwigg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                            MCDUDATIES SCORE COMPUTER: SECURATION TIEFS SCORE COMPUTER: BEASTER VERSION 1.5

CURENTARE: FASISTER VERSION 1.5

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,634

FILING DATE: 05-MAY-1995

CLASSIFICATION: 514

PRIOR APPLICATION NUMBER: 08/390,850

FILING DATE: FEBRUARY 17, 1995

APPLICATION NUMBER: 08/354,920

FILING DATE: December 13, 1994

APPLICATION NUMBER: 08/152,487

FILING DATE: December 13, 1994

APPLICATION NUMBER: 08/152,487

FILING DATE: No. 5731295ember 12, 1993

APPLICATION NUMBER: 17, 1992

ATTORNEY/AGENT INFORMATION:
NAME: WALDERY BICHARD

REFERENCE/DOCKET NUMBER: 23,327

REGISTRATION NUMBER: 23,327

REFERENCE/DOCKET NUMBER: 211/084

TELLERAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 590:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   364 TICITGGGGGCCCAGCT 380
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Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 90071
COMPUTER READABLE FORM:
                                                                                STREET: Suite 4700 CITY: Los Angeles STATE: California COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A.
                                                                                                                                                                                                                                            90071
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82.4%; Pred. No. 2.1e+02;
ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 323
US-08-435-634-590/c
| Sequence 590 Application US/08435634
| Patent NO. 5731295
| GENERAL INFORMATION:
| APPLICANT: Draper, Kenneth G. APPLICANT: PavCo, Pamela APPLICANT: McSwiggen, James APPLICANT: Gestioncomb, Dan T. TITLE OF INVENTION: METHOD AND REAGENT FOR TREATMENT; TITLE OF INVENTION: OF ARTHRITIC CONDITIONS | NUMBER OF SEQUENCES: 1151
| CORRESPONDENCE ADDRESS:
    APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Gustoffen, John
APPLICANT: Gustoffen, John
APPLICANT: Gustoffen, John
TITLE OF INVENTION: METHOD AND REAGENT FOR TREATMENT
TITLE OF INVENTION: OF ATTHRITIC CONDITIONS
NUMBER OF SEQUENCES: 1151
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon
STREET: 631 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM;
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: ENCRAGE
COMPUTER: ENCRAGE
COMPUTER: ENCRAGE
COMPUTER: ENGRAGE
COMPUTER: ENGRAGE
COMPUTER: ENGRAGE
COMPUTER: FASTER: TBM P.C. DOS 5.0
SOFTWARE: FASTER: TBM P.C. DOS 5.0
SOFTWARE: FASTER: TBM P.C. DOS 5.0
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/435,634
FILING DATE: FEDTUARY 17, 1995
FILING DATE: December 13, 1994
APPLICATION NUMBER: 08/354,920
FILING DATE: December 13, 1994
APPLICATION NUMBER: 08/354,920
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/989,848
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/989,848
FILING DATE: December 7, 1992
APPLICATION NUMBER: 211/084
FILING DATE: DECEMBER: 211/084
FILING DATE: DECEMBER: 211/084
FILING DATE: DECEMBER: 23.37
REFERENCE/DOCKET NUMBER: 32.327
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Best Local Similarity 82.43
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 17 base pairs
TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
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ANTI-SENSE: NO
ORIGINAL SOURCE:
INDIVIDUAL ISOLATE: Primer PGEX-R
US-08-466-033-104
    US 08/389,886
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GACCGTCTCCGGGAGCT
                        FILING DATE: 15-FEB-1
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA HYPOTHERT
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ZIP: 94306
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US-08-444-733-104
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 104, Application US/08466033

Patent No. 5766840

GENERAL INFORMATION:

APPLICANT: Kim, Jungauh P.

APPLICANT: Young, LaVonne M.

APPLICANT: Young, LaVonne M.

APPLICANT: Linnen, Jeffrey M.

TITLE OF INVENTION: Hepatitis G Virus and Molecular

TITLE OF INVENTION: Cloning Thereof

NUMBER OF SEQUENCES: 277

CORRESPONDENCE ADDRESS:

ADDRESSEE: Dehlinger & Associates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94306

ZIP: 94306

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,033
      MEDIUM TYPE: 5.0

MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
COMPRISE: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: IBM COMPATIBLE
SOFTWARE: FastSEQ Version 1.5

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,634
FILING DATE: 05-MAY-1995
FILING DATE: Peruary 17, 1995
APPLICATION NUMBER: 08/354,920
FILING DATE: December 13, 1994
APPLICATION NUMBER: 08/152,467
FILING DATE: NO. 5731295ember 12, 1993
APPLICATION NUMBER: 07/899,848
FILING DATE: NO. 5731295ember 12, 1993
APPLICATION NUMBER: 07/899,848
FILING DATE: NO. 5731295ember 12, 1993
APPLICATION NUMBER: 37,1992
ATORNEY/AGENT INFORMATION:
NAME: WALDLEY, RICHARC
REGISTRATION NUMBER: 32,327
REFERENCE/POCKET NUMBER: 211/084
TELLEPHONE: (213) 955-0440
TELLEPHONE: (213) 955-0440
TELLER: 67-3510
TELLEPHONE: APPLICATION: S92:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ....urkSSEE: Dehlinger & Associates
STREET: 350 Cambridge Ave., Suite 250
CITY: Palo Alto
COINTER: CA
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1056 CCCTGGCCTTCCCATCA 1072
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Best Local Similarity 58.8
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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US-08-435-634-592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 325
US-08-466-033-104
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Patent No. 5824507

GENERAL INFORMATION:

APPLICANT: Wages, John

APPLICANT: Young LaVonne M.

APPLICANT: Innen, Jeffrey M.

APPLICANT: Linnen, Jeffrey M.

TITLE OF INVENTION: Hepatitis G Virus and Molecular TITLE OF INVENTION: Cloning Thereof

NUMBER OF SEQUENCES: 277
PRICK APPLICATION NUMBER: US 08/357,509
FILING DATE: 16-DEC-1994
PRICK APPLICATION NUMBER: US 08/329,729
PRICK APPLICATION NUMBER: US 08/329,729
FILING DATE: 26-OCT-1994
PRICK APPLICATION NUMBER: US 08/344,271
PRICK APPLICATION NUMBER: US 08/245,558
FILING DATE: 23-NOV-1994
PRICK APPLICATION NUMBER: US 08/285,558
FILING DATE: 03-NUG-1994
PRICK APPLICATION NUMBER: US 08/285,543
FILING DATE: 03-NUG-1994
PRICK APPLICATION NUMBER: US 08/26,985
FILING DATE: 20-NAX-1994
APPLICATION NUMBER: US 08/26,985
FILING DATE: 20-NAX-1994
ATORNEY AGENT INFORMATION:
NAME: Fabian, Gary R.
FELENANIO NUMBER: 33,875
REGISTRATION NUMBER: 4600-0201.36/G100P11
TELECOMMUNICATION INPORMATION:
TELECOMM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Associates
STREET: 350 Cambridge Ave., Suite 250
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Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3; Indels
CCMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/710,134
FILING DATE: 13-SEP-1996
CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: Dugan, Deborah A.
REGISTRATION NUMBER: 137,315
REFERENCE/DOCKET NUMBER: 137,315
REFERENCE/DOCKET NUMBER: 165-8.1
TELECHMUNICATION INFORMATION:
TELECHMUNICATION INFORMATION:
TELECHMUNICATION INFORMATION:
TELECHMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

0.9%; Score 12.2; DB 1;
Best Local Similarity 82.4%; Pred. No. 2.1e+02;
Matches 14; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: August 17, 1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
INCluding application
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/008,895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: Dan B McSwiggen
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: BISEASES OR CONDITIONS
TITLE OF INVENTION: BISEASES OR CONDITIONS
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (1-CAM-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "Oligonucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERAL
SOFTWARE: WORD FELL.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-292-620A-1644/c
Sequence 1644, Application US/08292620A
Patent No. 5837542
GENERAL INFORMATION:
APPLICANT: Susan Grimm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: RIBOZYME TREAT TITLE OF INVENTION: DISEASES OR CONTITLE OF INVENTION: RELATED TO LIVITLE OF INVENTION: INTRACELLULAR TITLE OF INVENTION: MOLECULE-1 (1-NUMBER OF SEQUENCES: 2390 CORRESPONDENCE ADDRESS: ADDRESSE: Lyon & Lyon STREET: 613 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07/989,849
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                                                                                                                                                                                                                                                                                                                                   TELEFAX: 508-872-5415
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: Sulte ...
CITY: Los Angeles
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APPLICANT: SHUBER, ANTHONY P.
TITLE OF INVENTION: HIGH THROUGHDUT SCREENING METHOD FOR
TITLE OF INVENTION: SEQUENCES OR GENETIC ALTERATIONS IN NUCLEIC ACIDS
NUMBER OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
0
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                                                                            APPLICATION NUMBER: US/US/444,/33
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/389,886
FILING DATE: 15-FEB-1995
FILING DATE: 15-FEB-1995
FILING DATE: 16-DEC-1994
PRICK APPLICATION NUMBER: US 08/357,509
FILING DATE: 16-DEC-1994
PRICK APPLICATION NUMBER: US 08/329,729
FILING DATE: 26-OCT-1994
PRICK APPLICATION DATA:
APPLICATION NUMBER: US 08/344,271
FILING DATE: 23-NOV-1994
PRICK APPLICATION DATA:
APPLICATION NUMBER: US 08/285,588
FILING DATE: 03-AUG-1994
PRICK APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/285,543
FILING DATE: 03-AUG-1994
PRICK APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/246,985
FILING DATE: 03-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: FABIAN, GATY R.
REGISTRATION NUMBER: 4600-0201.36/G100P11
TELECOMMUNICATION NUMBER: 4600-0201.36/G100P11
                  PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INDIVIDUAL ISOLATE: Primer PGEX-R
                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/444,733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genzyme Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 49, Application US/08710134
Patent No. 5834181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (415) 324-0880
TELEFAX: (415) 324-0860
INFORMATION FOR SEQ ID NO: 104:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   625 GACCAGCTCCAGGAGCT 641
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 327
US-08-710-134-49/C
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CLASSIFICATION: 435
FRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/008,895
FILING DATE: DATURATY 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard 0.
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 208/149
TELEPHONE: (213) 489-1600
TELEPRONEY. (213) 955-0440
                                                                                                                                                                                                                                                                                                                                             0.9%; Score 12.2; DB 1; 70.6%; Pred. No. 2.1e+02; tive 2; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 330

US-08-292-620A-1700/C

Sequence 1700, Application US/08292620A

PATENT NO. 5837542

GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwiggen
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: BISEASES OR CONDITIONS
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390

CORRESPONDENCE ADDRESS:
ADDRESSES:
ADDRES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Los Angeles
STATE: California
COUNTY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: STORYER
COMPUTER: IBM COMPATIBLE
COMPUTER: Word Perfect 5.1
CURREMY APPLICATION DATA:
APPLICATION UNMBER: US/08/22,620A
FILING DATE: ANGUST 17, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                              961 CAGGACTGACCCCTCAC 977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 CAGCAUTUACCCCUCAC 17
TELEFAX: (21,
TELEX: 67.3510
; INFORMATION FOR SEQ ID NO: 16
; SEQUENCE CHARACTERISTICS:
: LENGTH: 17 base pairs
"VPB: nucleic acid
                                                                                                                                                                                                                                                                                                                                       Query Match 0.95
Best Local Similarity 70.65
Matches 12, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                      ; TYPE: nucleic aci
; STRANDEDNESS: six
; TOPOLOGY: linear
US-08-292-620A-1697
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0.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 2.1e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           **Sequence 1697, Application US/08292620A; Sequence 1697, Application US/08292620A; Patent No. 5837542; GENERAL INFORMATION; APPLICANT: Susan Grimm APPLICANT: Ban T. Stinchcomb APPLICANT: Dan T. Stinchcomb APPLICANT: Sean Sullivan APPLICANT: Renneth G. Draper TITLE OF INVENTION: RIBOZYME TREATMENT OF TITLE OF INVENTION: NTRACELLULAR ADHESION TITLE OF INVENTION: INTRACELLULAR ADHESION TITLE OF INVENTION: INTRACELLULAR ADHESION TITLE OF INVENTION: INTRACELLULAR ADHESION NUMBER OF SEQUENCES: 2390

**CORRESPONDENCE ADDRESS: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700

CITY: Los Annon-
         FILING DATE: December 7, 1992
ATTORNEY AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELECOMUNICATION INFORMATION:
TELEFAX: (213) 489-1600
TELEFAX: (7-3510
TELEFAX: (7-3510
INFORMATION FOR SEQ ID NO: 1644:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         208/149
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        623 GGGACCAGCTCCAGGAG 639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAWE: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 206
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17 GCGACCAGGACCAGGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-292-620A-1644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-292-620A-1697
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Parent No. 5837542
GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
ITILE OF INVENTION: DISEASES OR CONDITIONS
ITILE OF INVENTION: DISEASES OR CONDITIONS
ITILE OF INVENTION: NOLECULE. (I-CAM-1)
NUMBER OF INVENTION: MOLECULE. (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Saite 4700
STREET: Saite 4700
                                                                                                                                                                                                                Sequence 1743, Application US/08292620A Patent No. 5837542
                 623 GGGACCAGCTCCAGGAG 639
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                                                              17 GCGACCAGGACCAGGAG 1
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TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.9%;
Best Local Similarity 82.4%;
Matches 14; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17 GCGACCAGGACCAGGAG 1
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                        RESULT 332
US-08-292-620A-1743/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 333
US-08-292-620A-1796/c
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US-08-292-620A-1743
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TELEX: 6
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                                                                             Query Match 0.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 2.1e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 2.1e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIAME: COLLIONIA

COUNTER: U.S.A.

ZIP: 90071-2066

COMPUTER READABLE FORM:

MEDIUM TYPE: Storage

COMPUTE: LBM COMPALIA

COMPUTE: LBM COMPALIA

OPERATING SYSTEM: LBM P.C.

SOFTWARE: Word Perfect 5.1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/22,620A

CLASSIFICATION DATA: (1994

CLASSIFICATION DATA: (435)

PRIOR APPLICATION DATA: (645)

PRIOR APPLICATION NUMBER: 08/008,895

FILING DATE: Jenuary 19, 1993

APPLICATION NUMBER: 07/989,849

FILING DATE: December 7, 1992

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                         US-08-29-620A-1707/c

Sequence 1707, Application US/08292620A

Sequence 1707, Application US/08292620A

GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dames McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: NELATED TO LEVELS OF
TITLE OF INVENTION: MOLECULLIAR ADHESION
TITLE OF INVEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Warburg, Richard J.
RECISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                 623 GGGACCAGCTCCAGGAG 639
                                                                                                                                                                                                                                                    17 GCGACCAGGACCAGGAG 1
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TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 170
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Ly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: linear
US-08-292-620A-1707
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Best Local Similarity
Matches 14; Conserva
; TOPOLOGY: li
US-08-292-620A-1700
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Ouery Match

Ouery Matches

Ouery Matc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620A
FILING DATE: August 17, 1994
CLASSIFICATION DATA: Gescribed below:
PRIOR APPLICATION DATA: Gescribed below:
PRIOR APPLICATION DATA: Gescribed below:
APPLICATION NUMBER: 08/08/895
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
APPLICATION NUMBER: 20/989,449
FILING DATE: December 7, 1992
APPLICATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELEFONNUMICATION INFORMATION:
TELEFONNUMICATION:

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JS-08-292-620A-1934/C

Sequence 1934, Application US/08292620A

Patent No. 5837542

GENERAL INFORMATION:

APPLICANT: Susan Grimm
APPLICANT: Susan Grimm
APPLICANT: Sean Sullivan

APPLICANT: Sean Sullivan

APPLICANT: Kenneth G. Draper

TITLE OF INVENTION: RIBGZYME TREATMENT OF

TITLE OF INVENTION: RELATED TO LEVELS OF

TITLE OF INVENTION: RELATED TO LEVELS OF

TITLE OF INVENTION: NUTRACELLULAR ADHESION

TITLE OF INVENTION: MOLECULE.1 (1-CAM-1)
         APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
CORRESPONDENCE: 2390
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               623 GGGACCAGCTCCAGGAG 639
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: BY
TITLE OF INVENTION: D:
TITLE OF INVENTION: RI
TITLE OF INVENTION: HI
TITLE OF INVENTION: M
NUMBER OF SEQUENCES: ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-292-620A-1873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ð
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATURG SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620A
FILING DATE: AUGUST 17, 1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
PRIOR APPLICATION DATA: described below:
PRIOR APPLICATION DATA: described below:
PRIOR APPLICATION DATA: 07/989,849
FILING DATE: December 7, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEROME (213) 489-1600
TELEROME (213) 955-0440
INFORMATION FOR SEQ ID NO: 1796:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 Dase pairs
                                                                                                                      APPLICANT: Susan Grimm
APPLICANT: Busan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: BISEAGES OR CONTITIONS
TITLE OF INVENTION: DISEAGES OR CONTITIONS
TITLE OF INVENTION: MCLECTLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: LYON & LYON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: 801te 4700 CITY: Los Angeles STATE: California COUNTRY: U.S.A. ZIP: 90071-2066 COMPUTEN READABLE FORM: MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: Storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-292-620A-1873/c
Sequence 1873, Application US/08292620A
Fatent No. 583754.
GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dane T Stinchcomb
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
Sequence 1796, Application US/08292620A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     623 GGGACCAGCTCCAGGAG 639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17 GCGACCAGGACCAGGAG 1
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; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-292-620A-1796
                                     Patent No. 5837542
GENERAL INFORMATION:
APPLICANT: Susan (
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CORRESPONDENCE ADDRESS

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Length 17;
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APPLICANT: Wages, John
APPLICANT: Young, Lavonne M.
APPLICANT: Young, Lavonne M.
APPLICANT: Fry, Kirk E.
APPLICANT: Linnen, Jeffrey M.
TITLE OF INVENTION: Hepatitis G Virus and Molecular
TITLE OF INVENTION: Cloning Thereof
NUMBER OF SEQUENCES: 277
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Associates
STREET: 350 Cambridge Ave., Suite 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 94306

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATR:
APPLICATION NUMBER: US/08/464,134
                                                                                                                                                                                                                                                                                                                                                                                                                                              0.9%; Score 12.2; DB 1;
82.4%; Pred. No. 2.1e+02;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "Oligonucleotides"
                   CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Dugan, Deborah A.
REGISTRATION NUMBER: 37,315
REFERENCE/DOCKET NUMBER: GEN4-12.1
TELECOMMUNICATION:
TELEPHONE: 508-872-8400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AFLILING DATE:

CLASSIFICATION NOMBER: US/08/464,134
FILING DATE:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/359,886
FILING DATE: 15-FBB-1995
FILING DATE: 16-DEC-1994
FILING DATE: 16-DEC-1994
FILING DATE: 16-DEC-1994
FILING DATE: 26-OCT-1994
FILING DATE: 26-OCT-1994
FILING DATE: 26-OCT-1994
FRIOR APPLICATION NUMBER: US 08/329,729
FILING DATE: 23-NOV-1994
FRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/344,271
FILING DATE: 23-NOV-1994
FRIOR APPLICATION DATA:
APPLICATION UNDERF: US 08/344,271
FILING DATE: 03-AUG-1994
FRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 104, Application US/08464134
Patent No. 5849532
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        441 AAAGTTGCTGAAGTTTG 457
07-JUN-1995
                                                                                                                                                                             TELEFAX: 508-872-5415
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 AAATTTGATGAAGTATG
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.9%
Best Local Similarity 82.4%
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 350 Cambridge: STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 337
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Patent No. 5849483
GENERAL INFORMATION:
APPLICANT: SHUBER, ANTHONY P.
TITLE OF INVENTION: SEQUENCES OR GENETIC ALTERATIONS IN NUCLEIC ACIDS NUMBER OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
ADDRESSE: GENEYAL COPPORATION: COMPONDENCE ADDRESSE: GENEYAL ON THE ADDRESSE: GENEYAL OF THE ADDRESSE: GENEYAL COPPORATION
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                                                          CITY: Los Angeles
CITY: Los Angeles
COUNTRY: U.S.A.
ZIP: 30071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/22,620A
FILING DATE: August 17, 1994
CLASSIFICATION: 435
FILING DATE: MORE CONSUMENTIAL ON NUMBER: 08/08/895
FILING DATE: December 7, 1993
APPLICATION NUMBER: 08/008,895
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE DOCKET NUMBER: 208/149
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER REALGABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
GURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%; Score 12.2; DB 1;
82.4%; Pred. No. 2.1e+02;
tive 0; Mismatches 3;
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       623 GGGACCAGCTCCAGGAG 639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17 GCGACCAGGACCAGGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: One Mountain Road
CITY: Framingham
STATE: Massachusetts
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 82.49
Matches 14, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: 1
US-08-292-620A-1934
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APPLICATION NUMBER: US 08/344,271
FILING DATE: 23-NOV-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/285,558
FILING DATE: 03-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/285,543
FILING DATE: 03-AUG-1994
PRIOR APPLICATION NUMBER: US 08/246,985
FILING DATE: 20-NAY-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; ANTI-SENSE: NO
; ORIGINAL SOURCE:
; INDIVIDUAL ISOLATE: Primer PGEX-R
US-08-461-361-104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/389,886
FILING DATE: 15-FEB-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94306
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 625 GACCAGCTCCAGGAGCT 641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GACCGTCTCCGGGAGCT 17
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Best Local Similarity 82.45
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 17 base par
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     à
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Kim, Jungsuh P.
APPLICANT: Kim, Jungsuh P.
APPLICANT: Young, LaVonne M.
APPLICANT: Young, LaVonne M.
APPLICANT: Young, LaVonne M.
APPLICANT: Linnen, Jeffrey M.
TITLE OF INVENTION: Hepatitis G Virus and Molecular
TITLE OF INVENTION: Cloning Thereof
NUMBER OF SEQUENCES: 277
CORRESPONDENCE AUDRESS:
ADDRESSEE: Deblinger & Associates
STREET: 350 Cambridge Ave., Suite 250
CITY: Palo Alto
APPLICATION NUMBER: US 08/285,543
FILING DATE: 03-Aug-1994
PRIOR PAPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/246,985
FILING DATE: 20-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Fabian, Gary R.
REGISTRATION NUMBER: 33,875
REFRENCE/DOCKET WIMBER: 4600-0201.36/G100P11
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION INFORMATION:
TELECOMMUNICATION INFORMATION INFORMATION INFORMATION INFORMATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATE:
APPLICATION DATE:
FILING DATE: 15-FEB-1995
PRIOR APPLICATION DATE: 08/387,509
FILING DATE: 16-DEC-1994
PRIOR APPLICATION DATE: 16-DEC-1994
APPLICATION NUMBER: US 08/329,729
FILING DATE: 26-OCT-1994
FILING DATE: 26-OCT-1994
FILING DATE: 26-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: DNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
INDIVIDUAL ISOLATE: Primer PGEX-R
US-08-464-134-104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 104, Application US/08461361; Patent No. 5856134; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       625 GACCAGCTCCAGGAGCT 641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 GACCGTCTCCGGGAGCT 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 338
US-08-461-361-104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     qq
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0, Length 17; Indels Sequence 104, Application US/08485910

APPLICANT: Wages, John

APPLICANT: Young, LaVonne M.

APPLICANT: Linnen, Jeffrey M.

TITLE OF INVENTION: Hepatitis G Virus and Molecular ITILE OF INVENTION: Cloning Thereof

NUMBER OF SEGUENCES: 277

MUMBER OF SEGUENCES: 277

ADDRESSEE: Deblinger & Associates

STREET: 350 Cambridge Ave., Suite 250

CITY: Palo Alto NAME: Fabian, Gary R.
REGISTRATION NUMBER: 33,875
REGISTRATION NUMBER: 4600-0201.36/G100P11
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0880
TELEPHONE: (415) 324-0860
INFORMATION FOR SEQ ID NO: 104:
SEQUENCE CHARACTERISTICS: SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/485,910 0.9%; Score 12.2; DB 1; 82.4%; Pred. No. 2.1e+02; tive 0; Mismatches 3;

Thu Jan

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Retent No. 5882856

GENERAL INFORMATION:
APPLICANT: SHUBER, ANTHONY P.
TITLE OF INVENTION: UNIVERSAL PRIMER SEQUENCE FOR MULTIPLEX
ITTLE OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
ADDRESSEE: RAE-VENTER LAW GROUP
STREET: 2.50. Sheridan Ave., Ste. 440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 2.18+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                             PAPPLICATION NUMBER: US 08/344,271

PILING DATE: 23-NOV-1994

PRIOR APPLICATION NUMBER: US 08/285,558

PRIOR APPLICATION DATA:

PAPPLICATION NUMBER: US 08/285,558

PRIOR APPLICATION NATA:

PRIOR APPLICATION NATA:

PRIOR APPLICATION NATA:

PRIOR APPLICATION NUMBER: US 08/246,985

PRIOR APPLICATION NUMBER: US 08/240,086

PRI
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MEDIUM TYPE: Floppy disk
COMPUTER ISH PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
APPLICATION NUMBER: US/08/474,450A
FILING PATE: 7-JUNE-1995
           PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/329,729
FILING DATE: 26-0CT-1994
PRIOR APPLICATION DATA:
TOTAL TO NUMBER: US 08/344,271
APPLICATION NUMBER: US 08/357,509
FILING DATE: 16-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORIGINAL SOURCE:
; INDIVIDUAL ISOLATE: Primer PGEX-R
US-08-485-910-104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       625 GACCAGCTCCAGGAGCT 641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GACCGTCTCCGGGAGCT 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.9
Best Local Similarity 82.4
Matches 14, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
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STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: lines
MOLECULE TYPE: DI
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 340
US-08-474-450A-62
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GENERAL INFORMATION:
APPLICANT: MUBLIER, Rolf
APPLICANT: ZWICKER, Joerk
APPLICANT: ZWICKER, JOERK
APPLICANT: SEDIACEK, Hans-Herald
TITLE OF INVENTION: CYCLE-REGULATED EXPRESSION OF GENES AND THERAPEUTIC
TITLE OF INVENTION: METHODS OF UTILIZING SUCH CONSTRUCTS
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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0
                                                                                                                                                                                                                                                                                                                                          Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 12.2; DB 1; Length 17; Pred. No. 2.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: D.C.
CUUNTRY: USA
ZIONTRY: USA
ZIONTRY: USA
ZIONTRY: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: TEM PC compatible
COMPUTER: TEM PC compatible
COMPUTER: Tam PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/798,738
FILING DATE: 13-FEB-1997
CLASSIFICATION NUMBER: DE 1965274.2
FILING DATE: 13-FEB-1966
ATTONNEY/AGENT INFORMATION:
NAME: GENNADOS, PATRICIA D.
REGISTRATION NUMBER: 33,683
REFERENCE/DOCKET NUMBER: 33,683
REFERENCE/DOCKET NUMBER: 18748/318
                                                                                                                                                                                                                                       MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "WT-1 PRIMER SEQUENCE - N"
                                                                                                                                                                                                                                                                                                                                 0.9%; Score 12.2; DB 1;
82.4%; Pred. No. 2.1e+02;
tive 0; Mismatches 3;
REGISTRATION NUMBER: 32,750
REFERENCE/DOCKET NUMBER, GECO.001.00US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 328-4477
INFORMATION FOR SEQ ID NO: 62:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREE: Foley & Lardner STREE: 3000 K Street, N.W., Suite 500 STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 10, Application US/08798738 Patent No. 5885833
                                                                                                                                                                                                                                                                                                                                                                                                                     386 CAGAGGTGGCAGCAATG 402
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TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 82.4%
....hes 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPAX: (202)672-5399
TELEX: 904136
INFORMATION FOR SEQ ID NO;
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                       US-08-474-450A-62
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0.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 2.1e+02; Live 0; Mismatches 3; Indels
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APPLICANT: ARLEAT, Saghir
APPLICANT: Rell, Patricia
APPLICANT: McSwiggen, James
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT
TITLE OF INVENTION: OF DISEASES OR CONDITIONS RELATED
TITLE OF INVENTION: TO LEVELS OF EPIDERMAL GROWTH
TITLE OF INVENTION: TO LEVELS OF EPIDERMAL GROWTH
TITLE OF INVENTION: TO LEVELS OF EPIDERMAL
CORRESPONDENCE ADDRESS:
ADDRESSEE: 1970 & LyON
STREET: 613 West Fifth Street
STREET: Suite 470
CITY: LOS Angeles
COLIMINARY
COLI
                                                                                                                COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/181,664
FLIING DATE: JANUARY 14, 1994
CLASSIFICATION NUMBER: J1678
ATTORNEY/AGENT INFORMATION:
NAME: WetherILY 17, Ph.D., John R.
RESTSTRENCE/DOCKET NUMBER: 31678
REFERENCE/DOCKET NUMBER: 31678
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPRAX: (619) 455-5110
INFORMATION FOR SEQ ID NO: 64:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
"TOTAL AND THE COMPATION OF 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: 3.5" DOS 5.0 SOFTWARE: FASTENG FOR Windows 2.0 CURRENT APPLICATION DATA: MS/08/985,162 FILING DATE: 04 December 1997 CLASSIFICATION: 514 PRIOR APPLICATION: 514 PRIOR APPLICATION DATA: APPLICATION NUMBER: 60/036,476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 85, Application US/08985162; Patent No. 6057156; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1039 GACTCTTCCCACGACAG 1055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17 Grererecedadada 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                              California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 14; Conserv
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COUNTRY: U.S.A.
ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: CDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-181-664-64
                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ð
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                              ò
                              Gaps
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US-08-484-661A-17

i Sequence 17, Application US/08484661A

i Sequence 17, Application US/08484661A

i Patent No. 6001645

i APPLICANT: SLATER, MICHAEL R.

APPLICANT: HARNITT, JAMES R.

APPLICANT: HARNITT, JAMES R.

APPLICANT: HARNITT, JAMES R.

APPLICANT: HARNITT, JAMES R.

APPLICANT: HOUNGWITH, MODIFIED THERMOPHILIC DNA POLYMERASES

TITLE OF INVENTION: FROM THERMOTOGA NEAPOLITANA

ITILE OF INVENTION: FROM THERMOTOGA NEAPOLITANA

NUMBER OF SECUENCES: 51

CORRESPONDENCE ADDRESS:

ADDRESSEE: MEDLEN & CARROLL

STREET: 220 MONTCOMENT STREET, SUITE 2200

STATE: CALIFORNIA

COUNTRY: UNITED STATES OF AMERICA

INTEL 94104

COMPUTER READASE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPATION

MEDIUM TYPE: Ploppy disk

COMPUTER: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/484,661A

FLING DATE OF THE OF-JUN-1995

CLASSIFICATION NUMBER: US/027

REFERENCE/DOCKET NUMBER: PRMG-01175

TELECHMONICATION INFORMATION:

REFERENCE/OFMUNICATION INFORMATION:

REFERENCE/OFMUNICATION INFORMATION:

TELECHMONIC (415) 705-8410

INFORMATION FOR SEQ ID NO: 177:

SEQUENCE CHARACTERISTICS:

LERGYHH: 17 Dase pairs

TYPE: MINISTERMENTER OF ANDERSE

TYPE: MARNIT OF ANDERSE

TYPE: MARNIT OF ANDERSE

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                    Indels
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Patent No. 6025127
GENERAL INFORMATION:
APPLICANT: Sidransky, David
TITLE OF INVENTION: NUCLEIC ACID MUTATION DETECTION IN
TITLE OF INVENTION: HISTOLOGIC TISSUE
NUMBER OF SEQUENCES: 82
CORRESPONDENCE ADDRESS:
ADDRESSEE: Spensley Horn Jubas & Lubitz
STREET: 1880 Century Park East, Suite 500
                    ب.
ن
          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1199 GACCITCACACCICCCC 1215
                                                                                                 598 ACCAGCCTGAAGCCTGA 614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GACCTTGACACGTCCTC 17
                                                                                                                                                                                        17 AGCAGCCTGAGTCCTGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-484-661A-17
Matches 14; Conservative
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EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-181-664-64/c
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0
                                                                                                                             0.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 2.1e+02; Live 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Actions Applicant APPLICANT: Actions APPLICANT: Fell, Particia APPLICANT: Fell, Particia APPLICANT: McSwiggen, James TITLE OF INVENTION: BNZYMATIC NUCLEIC ACID TREATMENT TITLE OF INVENTION: TO LEVELS OF EPIDERWAL GROWTH TITLE OF INVENTION: TO LEVELS OF EPIDERWAL GROWTH TITLE OF INVENTION: TO LEVELS OF EPIDERWAL GROWTH CORRESPONDENCE ADDRESSE: 1877

CORRESPONDENCE ALDRESS: ADDRESS: ADDRESSEE: 633 West Fifth Street
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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82.4%; Pred. No. 2.1e+02;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTY: 0.5.A.

ZIE: 90071-2066

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: PRESEQ for Windows 2.0
SOFTWARE: FastSEQ for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/985,162
CLASSIFICATION: 514
PRIOR APPLICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/036,476
FILING DATE: 31 JANUARY 1997
ATTORNEY,AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERRANCH DATE: 31,332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 237, Application US/08985162
Patent No. 6057156
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                 629 AGCICCAGGAGCICIGC 645
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       645 CATCCCCCAAGACCTGG 661
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TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17 CAGCCTTCAAGACCTGG
                                                                                                                      Query Match
Best Local Similarity 82.49
Matches 14, Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 82.4
Matches 14; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                               ; TOPOLOGY: linear
US-08-985-162-104
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-985-162-237
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US-08-985-162-293
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TELEX: 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Aktuar, Saghir
APPLICANT: Rell, Parricia
APPLICANT: Rell, Parricia
APPLICANT: Rell, Parricia
APPLICANT: McSwiggen, James
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT
TITLE OF INVENTION: TO LEVELS OF EPDERMAL GROWTH
TITLE OF INVENTION: TO LEVELS OF EPDERMAL GROWTH
TITLE OF INVENTION: FACTOR RECEPTORS
NUMBER OF SEQUENCES: 1877
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: 8.0 Lorage
COMPUTER: 1EM COMPACIBLE
OPERATING SYSTEM: 1EM P.C. DOS 5.0
SOFTWARE: FASTSON FOR WINDOWS 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/985,162
FILING DATE: 04 December 1997
CLASSIFTATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/036,476
FILING DATE: 31 January 1997
APTONENY AGENT INFORMATION:
NAME: WAINLY RECEIVED NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REJECHONE: (213) 489-1600
TELLECHONE: (213) 499-1600
FILING DATE: 31 January 1997
ATTORNEY/AGENT INFORMATION:
NAME: WARDING, RICHARD J.
REGISTRATION NUMBER: 23.0.107
REFERENCE/DOCKET NUMBER: 230/107
TELECOMONIOTATION INFORMATION:
TELECHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: (213) 955-0440
TELEFAX: (713) 955-
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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STRANDEDNESS:
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US-08-985-162-104/C
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TOPOLOGY:
US-08-985-162-85
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APPLICANT: JARVIS, THALE
APPLICANT: MCSWIGGEN, JAMES A.
APPLICANT: MCSWIGGEN, JAMES A.
APPLICANT: MICHOGEN DAN T.
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES
TITLE OF INVENTION: OR CONDITIONS RELATED TO LEVELS OF C-FOS
FILE REPERENCE: 23.1/175
CURRENT APPLICATION NUMBER: US/08/998,099A
CURRENT FILING DATE: 1997-12-24
MARLIER PAPLICATION NUMBER: 08/37,658
EARLIER PAPLICATION NUMBER: 08/373,124
EARLIER PILING DATE: 1995-01-33
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                                                                                                                CONDITY: ULLILOW
ZIP: 94104
COMPUTER READABLE FORM:
MEDIUM TYEE: Floppy disk
CCMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
SOFTWARE: Patentin Release #1.0, Version #1.30
CURSENT APPLICATION DATA:
APPLICATION NUMBER: US/08/656,664
FILING DATE: 31-MAY-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAMME: Ingolia, Diane E.
REGISTRATION NUMBER: PRMG-02185
TELECOMMUNICATION INFORMATION:
TELEFAK: (415) 705-8410
TELEFAK: (415) 705-8410
TELEFAK: (415) 707-8338
INFORMATION FOR SEQ ID NO: 17:
                          220 Montgomery Street, Suite 2200
                                                                                      California
United States Of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 82, Application US/08998099A Patent No. 6103890
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MOLECULE TYPE: DNA (genomic)
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14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                        San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 11, Conserv
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US-08-998-099-82
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                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-998-099-82
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0
Sequence 293, Application US/06985162
Batent No. 6657166
GENERAL INPERMATION:
APPLICANT: Achtar, Saghir
APPLICANT: Achtar, Saghir
APPLICANT: Rell, Patricia
APPLICANT: Rell, Patricia
APPLICANT: Rell, Patricia
APPLICANT: RESWIGGEN, James
ITILE OF INVENTION: ENZYMATIC RELATED
TITLE OF INVENTION: TO LEVELS OF EPIDERMAL GROWTH
TITLE OF INVENTION: FACTOR RECEPTORS
NUMBER OF SEQUENCES: 1877
CORRESPONDENCE ADDRESS:
ADDRESSER: Lyon & Lyon
STREET: Gailte 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90701-2066
COMPUTER READALE FORM:
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
COMPUTER: Bateseg for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/985,162
FILING DATE: 04 December 1997
CLASSIFICATION DATA:
APPLICATION NUMBER: 05.327
REFERENCE/DOCKET NUMBER: 23.327
REFERENCE/DOCKET NUMBER: 23.327
TELECOMMUTICATION IN A AGA-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%; Score 12.2; DB 1; Length 17; 70.6%; Pred. No. 2.1e+02; tive 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: THERMOPHILIC DNA POLYMERASES FROM TITLE OF INVENTION: THERMOTOGA NEAPOLITANA NUMBER OF SEQUENCES: 57
CORRESPONDENCE ALDRESS:
ADDRESSEE: Medlen & Carroll
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17, Application US/08656664
| Patent No. 6077664
| GENERAL INFORMATION:
| APPLICANT: Slater, Michael R. APPLICANT: Huang, Fen
| APPLICANT: Hartnett, James R. APPLICANT: Bolchakova, Elena
| APPLICANT: Stortes, Douglas R. APPLICANT: Cotto, Paul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 293:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       577 CAGGCCCTCCGTCTGCC 593
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 70.6
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
TOPOLOGY:
US-08-985-162-293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 348
US-08-656-664-17
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                                                          APPLICAN:
APPLICAN:
TITLE OF INVENTION: RIBOZYRE .
TITLE OF INVENTION: DISEASES OR CONDITATION OF TITLE OF INVENTION: RELATED TO LEVELS OF TITLE OF INVENTION: INTRACELLULAR ADHESION TITLE OF INVENTION: MOLECULE-1 (I-CAM-1) NUMBER OF SEQUENCES: 2390
CORRESPONDENCES .2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Saile 4700
STREET: Saile 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1700, Application US/09071845
; Patent No. 6132967
; GENERAL INPORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Renneth G. Draper
APPLICANT: Renneth G. Draper
TITLE OF INVENTION: DISEASES OR CONDITIONS
                                                                                                                                                                                                                                                                                                                                                                                                               STREET: Solice 4700
CITY: Los Angeles
STREET: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
COMPUTER: MACOMPATIBLE FORM:
MEDIUM TYPE: GLORGE
COMPUTER: LAW
MEDIUM TYPE: GLORGE
COMPUTER: MACOMPATIBLE
COMPUTER: MACOMPATIBLE
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: AUGUST 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (213) 489-1600
TELERAX: (213) 955-0440
TELER: 67-3510
INFORMATION FOR SEQ ID NO: 1697:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  961 CAGGACTGACCCCTCAC 977
Dan T. Stinchcomb
James McSwiggen
Sean Sullivan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 CAGCAUUDACCCCUCAC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: US-09-071-845-1697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 352
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0.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 2.1e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                           Sequence:
Patent No. 613296,
Patent No. 613296,
GENERAL INFORMATION:
APPLICANT: Busan Grimm
APPLICANT: James McSwiggen
APPLICANT: Games McSwiggen
APPLICANT: Kenneth G. Draper
APPLICANT: Reneth G. Draper
The Condition
APPLICANT: Tenneth G. Draper
The Condition
The Condition of t
                                                                                                                                                                                                                                                                                                                                  APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: BLEATED TO LEVELS OF
TITLE OF INVENTION: NOLECULE. (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: LYON & LYON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 8torage
COMPUTER: IBM Compatible
COMPUTER: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REGISTRATION INFORMATION:
TELEPHONE: (213) 955-0440
TELEPAX: (213) 955-0440
TELERAX: 67-3510
TELERAX: 67-3510
SEQUENCE CHARACTERISTICS:
TENDORMATION FOR SEQ ID NO: 1644:
                                                                                       Sequence 1644, Application US/09071845
Patent No. 6132967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     623 GGGACCAGCTCCAGGAG 639
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nucleic acid
DEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                         RESULT 350
US-09-071-845-1644/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-071-845-1644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
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; Sequence 1697, Application US/09071845 ; Patent No. 6132967 ; GENERAL INFORMATION:

RESULT 351 US-09-071-845-1697

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0.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 2.1e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 354
US-09-071-845-1743/C
US-09-071-845-1743/C
Sequence 1743, Application US/09071845
Fatent No. 6132967
GERERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwigen
APPLICANT: James McSwigen
APPLICANT: Asea Sullivan
APPLICANT: Renacth G. Draper
ITILE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: RIATED TO LEVELS OF
TITLE OF INVENTION: NOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCES: 2390
                                                                                                    ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" DISKETTE, 1.44 Mb
MEDIUM TYPE: 3.5" DISKETTE, 1.00 COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
                                                                                                                                                                                                                                                                                                                      FILING DATE:

FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: OBLOB,895
FILING DATE: December 7, 1993
APPLICATION NUMBER: 07/89,849
FILING DATE: December 7, 1992
ATTORNEY, AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
FILEPHONE: (213) 489-1600
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
TELERX: 67-3510
INFORMATION FOR SEQ ID NO: 1707:
SEQUENCE CHARACTERISTICS:
LENGYTH: 17 Dase pairs
TYPE: nucleic acid
STRANDEDNESS: SINGLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   623 GGGACCAGCTCCAGGAG 639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17 GCGACCAGGACCAGGAG 1
             Suite 4700
                                     Los Angeles
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: Californi
COUNTRY: U.S.A.
ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
US-09-071-845-1707
                                   CITY: LOE
STATE: Ca
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 353
US-09-071-845-1707/c

| Sequence 1707, Application US/09071845
| Patent No. 6132967
| GENERAL INFORMATION:
| APPLICANT: Susan Grimm
| APPLICANT: Susan Grimm
| APPLICANT: Sean Sullivan
| APPLICANT: Sean Sullivan
| APPLICANT: Sean Sullivan
| TILE OF INVENTION: RIBOZYME TREATMENT OF
| TITLE OF INVENTION: BIBOZYME TREATMENT OF
| TITLE OF INVENTION: BIBAZES OR CONDITIONS
| TITLE OF INVENTION: MOLECULULAR ADHESION
| TITLE OF SEQUENCES: 2390
     RELATED TO LEVELS OF INTRACELLULAR ADHESION MOLECULE-1 (I-CAM-1): 2390
                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE:
CLASSIFICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/08,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,649
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TITLE OF INVENTION: RELATED TO LET TITLE OF INVENTION: INTRACELLULAR TITLE OF INVENTION: INTRACELLULAR NUMBER OF SEQUENCES: 2390.
CORRESPONDENCE ADDRESS: ADDRESS: ADDRESS: ADDRESSE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Shite 4700 CITY: Los Angeles STREET: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 499-1600
TELEXX: (213) 955-0440
TELEXX: 67-3510
INFORMATION FOR SEQ ID NO: 1700:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                i: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        623 GGGACCAGCTCCAGGAG 639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17 GCGACCAGGACCAGGAG
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ADDRESSEE: Lyon & Lyo
STREET: 633 West Fift
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                                                                                                                                                                                                                                           COUNTRY: U.S.A. ZIP: 90071-2066
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1873, Application US/09071845

Patent No. 6132967

GENERAL INFORMATION:
APPLICANT: Ban T. Stinchcomb
APPLICANT: James McSwiggen
APPLICANT: James McSwiggen
APPLICANT: Sans Sullivan
APPLICANT: Renneth G. Draper
ITILE OF INVENTION: RELATED TO LEVELS OF
ITILE OF INVENTION: RELATED TO LEVELS OF
ITILE OF INVENTION: RELATED TO LEVELS OF
ITILE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: 50146 4700
                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: US/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: WARDLEY, RECISTRATION NUMBER: 20,327
REGISTRATION NUMBER: 20,32,327
RESERENCE/DOCKET NUMBER: 20,327
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
INFORMATION FOR SEQ ID NO: 1796: SEQUENCE CHARACTERISTICS:
REPORT OF SEQ ID NO: 1796: SEQUENCE CHARACTERISTICS:
REDER OF SEQ ID NO: 1796: SEQUENCE CHARACTERISTICS:
REDER OF SEQ ID NO: 1796: SEQUENCE CHARACTERISTICS:
REDER OF SEQ ID NO: 1796: MEDER O
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ZIE: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
WEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: World Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US/08/292,620
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,6
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
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Best Local Similarity 82.45
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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US-09-071-845-1796/C
; Sequence 1796, Application US/09071845
; Fatent No. 6122367
; GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: James McSwiggen
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RIBOZYME ADHESION
TITLE OF INVENTION: INTRACELLUAR ADHESION
TITLE OF INVENTION: INTRACELLUAR ADHESION
TITLE OF INVENTION: NOISCULE-1 (I-CAM-1)
CORRESPONDENCE ADDRESSE:
ADDRESSEE: LYON & INVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A.

ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM COMPALIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE:
                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNDER: US/08/292,620
FILING DATE: AUGUST 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         208/149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/14
TELECOMUNICATION INFORMATION:
TELEPAN: (213) 955-0440
TELEPAN: (213) 955-0440
INFORMATION FOR SEQ ID NO: 1743:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700 CITY: Los Angeles STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           623 GGGACCAGCTCCAGGAG 639
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US-08-961-810-104/c
i Sequence 104, Application US/08961810
i Patent No. 6165713
i GENERAL INFORMATION:
APPLICANT: Liskay, Robert M.
APPLICANT: Baker, Sean M.
APPLICANT: Baker, Sean M.
APPLICANT: Bollag, Roni J.
APPLICANT: Rolodner, Richard D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS RELATING TO DNA
TITLE OF INVENTION: MISWATCH REPAIR GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 2.1e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                    0.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 2.1e+02;
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                                                                                                                                                                                                                                                                                     3: Indels
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Kolisch, Hartwell, Dickinson, McCormack
ADDRESSEE: Heuser
STREET: 520 S.W. Yamhill Street, Suite 200
CITY: Portland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: Oregon
COUNTRY: U.S.A.
ZIP: 97204
COMPUTER EADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/961,810
FILING DATE:
                                                                                                                                                                                                                                                                                     0; Mismatches
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ATTORNEY/AGENT INFORMATION:
NAME: Van Rysselberghe, Pierre C.
REGISTRATION NUMBER: 33,557
REFREENCE/DOCKET NUMBER: 04SU 306B
TELECHONICATION INFORMATION:
TELEPHONE: (503) 224-6655
TELEFAX: (503) 295-6679
            IDEDEX: 67-3510
INFORMATION FOR SEQ ID NO: 1934:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                   623 GGGACCAGCTCCAGGAG 639
                                                                                                                                                                                                                                                                                                                                                                                17 GCGACCAGGACCAGGAG 1
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TELEX: 360619
INFORMATION FOR SEQ ID NO: 3EQUENCE CHARACTERISTICS: LENGTH: 17 base pairs TYPE: nucleic acid STRANDEDNESS: single
  (213) 955-0440
                                                                                                                                                                                                                                    Query Match 0.9%
Best Local Similarity 82.4%
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
                                                                                   IENCTH: 17 Date TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
                                                                                                                                                                 ; TOPOLOGY: 1
US-09-071-845-1934
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: Dan T. Stinchcomb
APPLICANT: Sean SUllivan
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZDENE TREATMENT OF
TITLE OF INVENTION: RIBOZDENES OR CONDITIONS
TITLE OF INVENTION: RIBOZDENES OF
TITLE OF INVENTION: NURACELIULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: STORAGE
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PETFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
                                       NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REERENCE/DOCKET NUMBER: 208/149
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 1873:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANBENNES: single
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
FELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1934, Application US/09071845
Patent No. 6132957
GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: James McSwiggen
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                623 GGGACCAGCTCCAGGAG 639
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Best Local Similarity 82.4
Matches 14; Conservative
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California
                                                                                                                                                                                                                                                                                                                                 linear
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STATE: Califorr
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-071-845-1934/c
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                                                                                                RESULT 359
US-08-352-902D-104/c
; Sequence 104, Application US/08352902D
; Patent No. 6191268
; GENERAL INFORMATION:
; APPLICANT: Liskay, Robert M.
; Bronner, C. Eric
Baker, Sean M.
; Bollag, Roni J.
; Kolodner, Richard D.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS RELATING TO DNA
MISMATCH REPAIR GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: 1 OTHER INFORMATION: /note= "primers directed to genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTER: 0....

ZIP: 97204

COMPUTER READABLE FORM:

MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OMPUTER: IBM PC compatible
OMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/352,902D
FILING DATE: 09-Dec-1994
CLASSIFICATION: cUnknown>
ATTORNEY/AGRYT INFORMATION:
NAME: Van Rysselberghe, Pierre C.
REGISTRATION NUMBER: 33.557
REGISTRATION INFORMATION:
TELEPAX: (503) 295-6679

INFORMATION FOR SEQ ID NO: 104:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 2.1e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 520 S.W. Yamhill Street, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE DESCRIPTION: SEQ ID NO: 104:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 360
US-08-983.466-93
; Sequence 93, Application US/08983466
; Patent No. 6207372
; GENERAL INFORMATION:
; APPLICANT: SHUBER, ANTHONY P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         941 AGAGGTGTGAGGGCAGA 957
941 AGAGGTGTGAGCGCAGA 957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 AGACGTGAGAGCCCAGA 1
                                        17 AGACGTGAGAGCCCAGA 1
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STATE: Oregon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                intron DNA
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Gaps
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    UNIVERSAL PRIMER SEQUENCE FOR MULTIPLEX
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                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: TEM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN DATA:
APPLICATION NUMBER: US/08/983,466
FILING DATE: 10-FEB-1998
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: 08/44,450
FILING DATE: 07-JUNE-1995
PRIOR APPLICATION NUMBER: MO96/41012
APPLICATION NUMBER: MO96/41012
FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "WT-1 PRIMER SEQUENCE - N"
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US-09-091-590A-26
| US-09-091-590A-26
| Sequence 26, Application US/09091590A
| Patent No. 6242574
| GENERAL INFORMATION:
| APPLICANT: Nielsen, Klaus
| APPLICANT: Rroll Kristensen, Anne
| APPLICANT: Brunstedt, Janne
| TITLE OF UNVENTION: Anti-Microbial Proteins
| FILE REFERENCE: S-137-1101/MA/A/SGS/PCT |
| CURRENT APPLICATION NUMBER: US/09/091,590A |
| PRIOR PRILING DATE: 1999-05-06 |
| PRIOR FILING DATE: 1996-12-20 |
| PRIOR FILING DATE: 1996-12-21 |
| PRIOR FILING DATE: 1996-12-21 |
| PRIOR FILING DATE: 1996-12-21 |
| PRIOR FILING DATE: 1995-12-21 |
| PRIOR FILING 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Rae-Venter, Barbara
REGISTRATION NUMBER: 32,750
REFERENCE/DOCKET NUMBER: GECO.001.01US
TELECOMUNICATION INFORMATION:
TELEPHONE: (650) 328-4400
INFORMATION FOR SEQ ID NO: 93:
SEQUENCE CHARACTERISTICS:
                                                 DNA AMPLIFICATION 95
TITLE OF INVENTION: UNIVERSAL PRIM
TITLE OF INVENTION: DNA AMPLIFICAT
NUMBER OF SEQUENCES: 95
CORRESPONDENCE ADDRESS:
ADDRESSEE: RAE-VENTER LAW GROUP
STREET: 260 Sheridan Ave., Ste.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              386 CAGAGGTGGCAGCAATG 402
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                  CITY: Palo Alto
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 14; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                       94306
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SEQ ID NO 26
LENGTH: 17
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GENERAL INFORMATION:
APPLICANT: Shannon, Karen W.
APPLICANT: Shannon, Karen W.
APPLICANT: Belenstarr, Glenda C.
APPLICANT: Webb, Peter G.
APPLICANT: Webb, Peter G.
APPLICANT: Wicaid, Robert H.
ITILE OF INVENTION: Methods for evaluating oligonucleotide
ITILE OF INVENTION: probe sequences
NUMBER OF SEQUENCES: 1165
CORRESPONDENCE ADDRESS:
ADDRESSEE: Records Manager, Legal Department, Hewlett-Packard Company M/S 20
STREET: 3000 Hanover Street
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 2.1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/021,701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Cohen, Daniel
APPLICANT: Cohen, Daniel
APPLICANT: Blumenfeld, Marta
APPLICANT: Ilya, Chumakov
APPLICANT: Ilya, Chumakov
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: PROSTATE
FILE REFERENCE: GENSET.18CPICP
CURRENT PELLICATION NUMBER: US/09/338,907
CURRENT FILING DATE: 1999-06-23
EARLIER APPLICATION NUMBER: 08/996,306
FARLIER FILING DATE: 1997-12-22
EARLIER PILING DATE: 1997-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 UMBER: US/09/021,701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                     Sequence 111, Application US/09021701
Patent No. 6251588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 84, Application US/09338907
Patent No. 6265546
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 10-FEB-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Choi, Wendy A.
REGISTRATION NUMBER: 36,697
REPRESENCE/DOCKET NUMBER: 1097:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-236-2386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      296 TGTCTGCTGTGGGGGCT 312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 650-852-8063
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.9
Best Local Similarity 82.4
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-338-907-84/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                   94304
                                                JS-09-021-701-111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Records Manager, Legal Department, Hewlett-Packard Company M/S 20 STREET: 3000 Hanover Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ô
                                                                                                                                                                                                                                                                                                                                                                                           RESULT 362
US-09-021-701-53/c
; Sequence 53, Application US/09021701
; Patent No. 6251588
; GENERAL INFORMATION:
; APPLICANT: Shannon, Karen W.
; APPLICANT: Wolber, Paul K.
; APPLICANT: Webb, Peters G.
; APPLICANT: Kincaid, Robert H.
; TITLE OF INVENTION: Methods for evaluating oligonucleotide
; TITLE OF INVENTION: probe sequences
; TITLE OF UNDENTION: probe sequences
; NUMBER OF SEQUENCES: 1165
; CORRESPONDENCE ADDRESS: 1165
                                                                                                                                                                                                                                                  0
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                                                                                                                                                                                                 0.9%; Score 12.2; DB 1; Length 17; 64.7%; Pred. No. 2.1e+02; ive 3; Mismatches 3; Indels
                                                                                                                                                                                                                                           3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/021,701
FILING DATE: 10-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
ATTORNAY AGENT INFORMATION:
NAME: Choi, Wendy A.
REGISTRATION NUMBER: 36,697
REFERENCE/DOCKET VINBER: 10971464-1
TELECOMMUNICATION:
                NAME/KEY: misc_feature
LOCATION: (1)...(17)
OTHER INFORMATION: primer
LOCATION: (1)...(17)
OTHER INFORMATION: n = a, t, c, or g
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10-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1249 GCCATGTGAGGCCAGGT 1265
                                                                                                                                                                                                                                                                                         294 AATGICIGCIGIGGGG 310
                                                                                                                                                                                                                                                                                                                   1 AAYGINTGYTGYGCNGG 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 650-852-8063
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                   Best Local Similarity 64.7
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 14; Conserva'
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                                                                                                                                                  US-09-091-590A-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; ANTI-SENSE:
US-09-021-701-53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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FEATURE:
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0.9%; Score 12.2; DB 1; Length 17; 58.8%; Pred. No. 2.1e+02; tive 4; Mismatches 3; Indels
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Fatent No. 6346398
GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: Escobedo, Jaime
ITILE OF INVENTION: METHOD AND REAGENT FOR THE
ITILE OF INVENTION: TRAATWENT OF DISEASES OR
ITILE OF INVENTION: CONDITIONS RELATED TO LEVELS
ITILE OF INVENTION: GROWTH FACTOR
INTERPREDICTION: GROWTH FACTOR
ADDRESSEE: Lyon & Lyon
ADDRESSEE: Lyon & Lyon
    CONDITIONS RELATED TO LEVELS
OF VASCULAR ENDOTHELIAL
GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Btorage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRICA APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg Richard J.
REGISTRATION NUMBER: 218/064
TELECOMMUTCATION INFORMATION:
TELEPHONE: (213) 955-0440
TELEPHONE: (213) 955-0440
TELEPRA (7-35) 10 NO: 1909:
SEQUENCE CHARACTERISTICS:
TENGTH: TABER TO SEG ID NO: 1909:
                                                                                                                                                                                                                                                   ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
                                                                                                             ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             234 TCAGGCATCTGCATCTG 250
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                                                                          8502
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Matches 10; Conservative
                       TITLE OF INVENTION: OF TITLE OF INVENTION: GROW NUMBER OF SEQUENCES: 85CORRESSPONDENCE ADDRESS: ADDRESSE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                     CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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                                                                                                                                                                                                                                  COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-584-040-1922/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 2.1e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature

LOCATION: 1..17

CTHER INFORMATION: sequencing oligonucleotide moPGrace5R444

US-09-218-207-84
                                                                                                                                                                                                                           NAME/KEY: misc feature
i LOCATION: 1..17
CTHER INFORMATION: sequencing oligonucleotide moPGrace5R444
US-09-338-997-84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1909, Application US/08584040
Patent No. 6346398
GENERAL INFORMATION
APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Bscobedo, Jaine
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 84, Application US/09218207
Fatent No. 6346381
GENERAL INFORMATION:
APPLICANT: Cohen, Daniel
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Bougleleret, Lydie
TITLE OF INVENTION: Prostate cancer gene
FILE REFERENCE: GENSET.018CP1
CURRENT APPLICATION NUMBER: US/09/218,207
CURRENT APPLICATION NUMBER: 08/996,306
EARLIER FILING DATE: 1997-12-22
EARLIER APPLICATION NUMBER: 60/099,658
FARLIER PRILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 578
SOFTWARE: PATENT. DM
EARLIER FILING DATE: 1998-09-09
EARLIER APPLICATION NUMBER: 09/218,207
EARLIER FILING DATE: 1998-12-22
NUMBER OF SEQ ID NOS: 578
                                                                                                                                                                                                                                                                                                                                                                                                                                      253 ACCGACCTCCTGGGCTG 269
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Best Local Similarity 82.4-
Best Local 4, Conservative
                                                                                                                                                                                                                                                                                                                                                                                           14; Conservative
                                                                                                                                                             TYPE: DNA ORGANISM: Mus Musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Mus Musculus
                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
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US-09-218-207-84/c
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US-08-584-040-1909
                                                                                                             SEQ ID NO 84
LENGTH: 17
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LENGTH: 17
                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 2224 Application US/08584040
Fatent No. 6346398
GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: Bavco, Pamela
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZDF: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Accrage
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COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRICR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: WAIDLICY RICHARD J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECHONE: (213) 489-1600
TELECHONE: (213) 955-0440
TELEFAX: (213) 955-0440
TELEFAX: 67-3510
SEQUENCE CHARACTERISTICS:
TENGTH: 17 base pairs
                           FILING DATE: October 26, 1995
ATTOREY/AGENT INFORMATION:
NAME: Warburg, Richard J
REGIGSTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                       TELBEAX: (213) >>> TELBEAX: (213) >>> TELBEAX: 67-3510 |
| TELEX: 67-3510 |
| INFORMATION FOR SEQ ID NO: 2028: SEQUENCE CHARACTERISTICS: LENGTH: 17 base pairs | TYPE: nucleic acid | STRANDEDNESS: single | STRANDEDNESS: single |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          843 AGATGGGTCAGCATACC 859
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Best Local Similarity 64.77
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-584-040-2224/c
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US-08-584-040-2028
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-US-SHAUDO-2028
Sequence 2028, Application US/08584040
Fatent No. 6346398
GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: Bacobedo, James
APPLICANT: Escobedo, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STRATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READBABLE FORM:
MEDIUM TYPE: 315" Diskette, 1.44 Mb
MEDIUM TYPE: 315" Diskette, 1.44 Mb
COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage computed co
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OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
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CLASSIFICATION: 514
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.99
Best Local Similarity 82.49
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-584-040-1922
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US-08-584-040-2028
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nucleic acid

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Length 17;
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                                                                                                                                                                                                                                             APPLICANT: Parco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Stinchomb, Dan T
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: TREATHENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: GROWTH FACTOR
TITLE OF INVENTION: GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.9%; Score 12.2; DB 1;
58.8%; Pred. No. 2.1e+02;
tive 4; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. Darocon
GENERAL INCRMATION:
APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Storage
COMPUTER: IBM COMPACTÈDIE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: WAZDLOY, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUTCATION INFORMATION:
TELEPHONE: (213) 955-0440
TELEPHONE: 67-3510
TELEPACE: 67-3510
SEQUENCE CHARACTERISTICS:
TENDOMET'S BEED TO SEED TO S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                Sequence 3739, Application US/08584040
Patent No. 6346398
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 3840, Application US/08584040; Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1240 CTGGACGTGGCCATGTG 1256
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Best Local Similarity 58.55
Best Local Similarity
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: METATILLE OF INVENTION: CONITILE OF INVENTION: CONITILE OF INVENTION: OF VITTLE OF INVENTION: GROWNBER OF SEQUENCES: 85CORRESPONDENCE ADDRESSES: ADDRESSEE: Lyon & Lyor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-584-040-3739
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US-08-584-040-3840
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                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                     0;
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                                                                                                                                                            Query Match 0.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 2.1e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           METHOD AND REAGENT FOR THE TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL GROWTH PACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPB: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION TO STA
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY AGENT INFORMATION:
NAME: WARDLEY, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
RELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION INFORMATION:
TELECOMMUNICATION INFORMATION I
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ZIE: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 2554, Application US/08584040; Patent No. 6346398; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Pavco, Pamela
APPLICANT: Pavco, James
APPLICANT: Strinchconb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: METHOD AND JITLE OF INVENTION: TREATMENT OF ITLE OF INVENTION: CONDITIONS TITLE OF INVENTION: GROWTH PACTY
TITLE OF INVENTION: GROWTH PACTY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: Lyon & Lyon
633 West Fifth Street
Suite 4700
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Los Angeles
STATE: California
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                ; TOPOLOGY: 1
US-08-584-040-2224
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US-08-584-040-2554
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
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Gaps

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0.9%; Score 12.2; DB 1; Length 17; 23.5%; Pred. No. 2.16+02;
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Patent No. 6346398

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Barco, Pamela

APPLICANT: Barchchomb, Dan T.

APPLICANT: Stinchcomb, Dan T.

TITLE OF INVENTION: ONDITIONS RELATED TO LEVELS

TITLE OF INVENTION: GROWTH PACTOR

NUMBER OF SEQUENCES: 8502

CORRESPONDENCE ADDRESS:

ADDRESSE: Lyon & Lyon

STREET: Suite 4700

STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
                                                                        MEDIUM TIFE.

MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REGIOUS NUMBER: 32,327
REGIONALION NUMBER
             APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1142 CCTTTTTTTTTGG 1158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-584-040-3912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-584-040-3911
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 373
UG-08-564-040-3911
Sequence 3911, Application US/08584040
Patent No. 6346398
GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: MCSwiggen, James
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: CROWITH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCES: 8502
CORRESPONDENCES: 8502
CORRESPONDENCES: 1504
STREET: SAIM West Fifth Street
STREET: SAIM est Fifth Street
STREET: California
TREATMENT OF DISEASES OR
CONDITIONS RELATED TO LEVELS
OF VASCULAR ENDOTHELIAL
GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIPICATION: 514
PRIOR APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELECHONE: (213) 955-0440
TELECHONE: (213) 955-0440
TELECHONE: (713) 955-0440
TELECHONE: 67-3510
INFORMATION FOR SEQ ID NO: 3840:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
                        TITLE OF INVENTION: CONDITIONS REL
TITLE OF INVENTION: OF VASCULAR EN
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Suite 4700
CITY: Los Angeles
STREET: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1073 GGCAGGCTCTTCAGTGA 1089
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Best Local Similarity 58.8
Matches 10; Conservative
      INVENTION:
                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U.S.A.
ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-584-040-3840
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      TITLE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        qq
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                                                                                                                                                 Query Match 0.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 58.8%; Pred. No. 2.1e+02; Matches 10; Conservative 4; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:

APPLICANT: Pavco, Pamela
APPLICANT: Bavco, Pamela
APPLICANT: Stinchcomb, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: TREATHON AND REAGENT FOR THE
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: Suite 4700
CITT: LOS ANGELSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.9%; Score 12.2; DB 1; Best Local Similarity 82.4%; Pred. No. 2.1e+02; Matches 14; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Los Augeles
STATE: California
CONDATA: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: BISM Compatible
COMPUTER: WASTER: WAS PERCENTED
SOFTWARE: WASTER: TANG PERCENT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: OCTOBE: 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: WAEDLEY, RICHARG J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   218/064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REPRENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                          1073 GGCAGGCTCTTCAGTGA 1089
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                                                                                                                                                                                                                                                                                         1 GGCAUGGUCUUCUGUGA 17
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
          17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 17 base pairs
                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                linear
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                                                                                                        US-08-584-040-5441
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                                                                                   TOPOLOGY:
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O.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 29.4%; Pred. No. 2.1e+02;
Matches 5; Conservative 9; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Pavco, pamela
APPLICANT: Resigner, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Scobedo, Jaine
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATHENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELAIED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Los Angeles
COUNTY: Los Angeles
COUNTY: Los Angeles
COUNTY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD Perfect 5.1
CURRENT APPLICATION DATA:
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRICA APPLICATION: 514
PRICA APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY AGENT THYORMATION:
MANAN: MAANAL MAANAL
               FILING DATE: OCCODER 26, 1995
ATORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEFAX: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: (213) 955-0410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 5441, Application US/08584040 Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
APPLICATION NUMBER: 60/005,974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: LYON & LYON
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1146 TTTTTCTTTTGGAAGT 1162
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 UUGUUGCUUUUGGAAGU 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                          ;
US-08-584-040-3912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 375
US-08-584-040-5441
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0.9%; Score 12.2; DB 1; Length 17; 32.4%; Pred. No. 2.1e+02;
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Fatent No. 6346398
GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Dam T.
APPLICANT: Stinchomb, Dan T.
APPLICANT: Stinchomb, Dan T.
APPLICANT: Becobedo, Jaimes
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS
                       TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS
    METHOD AND REAGENT FOR THE
                                                           OF VASCULAR ENDOTHELIAL GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                          CITY: Los Angeles
STARE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
FILING DATE: January 11, 1996
CLASSIFICATION: 514
TITLE OF INVENTION: METHOD AND REF
TITLE OF INVENTION: TREATMENT OF L
TITLE OF INVENTION: OF VASCULAR EN
TITLE OF INVENTION: OF VASCULAR EN
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 6502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            878 CCAAGTTCCAGGAGCTG 894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17 ccaaggrcaaggrgcrg 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        82.48;
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INFORMATION FOR EGO ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ouery Match
Best Local Similarity 82.44
Matches 14, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 379
US-08-584-040-7406/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
US-08-584-040-7260
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                                                                                                                                                                          APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Beroched, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: PREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STRATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 MD
MEDIUM TYPE: 3.5" Diskette, 1.44 MD
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: October 26, 1995
ATTORNEY/AGBNT INFORMATION:
NAME: MAIDUTG, Richard J.
NAME: MAIDUTG, Richard J.
NAME: MAIDUTG, Richard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 378
US-08-584-040-7260/c
; Sequence 7260, Application US/08584040
; Patent No. 6346394
; GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
; APPLICANT: Stinchcom, James
; APPLICANT: Stinchcom, James
; APPLICANT: Stinchcom, James
                                                                                                             Sequence 5925, Application US/08584040 Patent No. 6346398 GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    218/064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 216
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 5925
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         881 AGTICCAGGAGCIGCGG 897
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       17 GTAGTCAGGAGCCCGCA
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Best Local Similarity 82.4
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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                                                                         RESULT 377
US-08-584-040-5925/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-584-040-5925
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 7877, Application US/08584040

Patent No. 646398

GENERAL INFORMATION:
APPLICANT: Paveo, Pamela
APPLICANT: Paveo, Pamela
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaimes
TITLE OF INVENTION: METATHOD AND REAGENT FOR THE
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
TITLE OF INVENTION: GROWTH FACTOR
CORRESPONDENCES: 8502
CORRESPONDENCES: 8502
ADDRESSED: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Sulte 4700
CITY. ICA ANDERSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPOTER: IBM COMPATIBLE
COMPOTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION 1514
PRICA APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: BLORAGE
PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,877
REFERENCE/DOCKET NUMBER: 32,877
REFERENCE/DOCKET NUMBER: 32,877
REFERENCE/DOCKET NUMBER: 32,876
TELEPAX: (213) 955-0440
TELEPAX: (213) 955-0440
TELEPAX: (7-3510
INFORMATION FOR SEQ ID NO: 7591:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STANDENDESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 7877:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   234 TCAGGCATCTGCATCTG 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 UCAAGCCUCUGCAUUUG 17
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(213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-584-040-7591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-584-040-7877
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                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 381
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 2.1e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            METHOD AND REAGENT FOR THE TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S.A.

ZIP: 90071-206
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: STORAGE
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PETFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
  COUNTRY: U.S.A.

ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM COMPALIAL
COMPUTER: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIGR APPLICATION: 514
FILING DATE: OCCODER 26, 1995
ATTONNEY ACENT THY MANDER: 60/005,974
FILING DATE: OCCODER 26, 1995
ATTONNEY ACENT THY MANDER: APPLICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 7551, Application US/08584040
Sequence 7551, Application US/08584040
Patent No. 6346398
GENERAL INFORMATION:
APPLICANT: Barco, Pamela
APPLICANT: Brinchoomb, Dan T.
APPLICANT: Stinchoomb, Dan T.
APPLICANT: Brinchoomb, Dan T.
APPLICANT: Brinchoomb, Dan T.
APPLICANT: Brinchoomb, Dan T.
APPLICANT: Broched, Jaime
TITLE OF INVENTION: METHOD AND REAGEN
TITLE OF INVENTION: CONDITIONS RELATE
TITLE OF INVENTION: GROWTH FACTOR
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET UNBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEFAX: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: (713) 955-0440
TELEFAX: (713) 955-0440
TELEFAX: (713) 955-0440
INFORMATION FOR SEQ ID NO: 7406:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       January 11, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1327 GTAGATCTTGTGTTTCA 1343
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-584-040-7406
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US-08-584-040-7591
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4.7%; Pred. No. 2.1e+02;
ve 3; Mismatches 3; Indels
Score 12.2; DB 1; Length 17;
Pred. No. 2.1e+02;
1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPOSITION AND METHODS FOR MODULATION OF GENE EXPRESSION IN PLANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: UJy 12, 1996
CLASSIFICATION NUMBER: 00/001,135
FILING DATE: UNLY 13, 1995
APPLICATION NUMBER: 60/001,135
FILING DATE: UNLY 13, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISCHMURICATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 319/247
TELECOMMUTICATION INPORMATION:
TELECOMMUTICATION INPORMATION:
                                                                                                                                                                                                                                                                                                  APPLICANT: Zwick, Michael G.
APPLICANT: Edington, Brent E.
APPLICANT: Edington, Brent E.
APPLICANT: Merlo, Patricia Ann Owens
APPLICANT: Guo, Lining
APPLICANT: Skokut, Thomas A.
APPLICANT: Young, Scott A.
APPLICANT: Folketts, Otto
APPLICANT: Folketts, Otto
TITLE OF INVENTION: COMPOSITION AND MET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Los Angeles
START: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READBLE FORM:
MEDIUM TYPE: Storage
                                                                                                                                                                                                                                     Sequence 216, Application US/08679645
Patent No. 6350934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 613 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             221 GAGCTCCTCAGCCTCAG 237
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                                                                                          939 GAAGAGGTGTGAGCGCA 955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 216
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.9%;
Best Local Similarity 64.7%;
Matches 11; Conservative
                                                                                                                                1 gaagaagucgaggga 17
      0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17 base pairs
      Query Match
Best Local Similarity 76.55
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-679-645-216
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                                                                                                                                                                                                 RESULT 383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        g
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                                                                                                                                                         Query Match

0.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 70.6%; Pred. No. 2.1e+02;
Matches 12; Conservative 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPOSITION AND METHODS FOR MODULATION OF GENE EXPRESSION IN PLANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTY: CALLLO-COUNTY: U.S.A.

ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: JULY 12, 1996
FILING DATE: JULY 12, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zwick, Michael G.
Edington, Brent E.
McSwiggen, James A.
Merlo, Patricia Ann Owens
Guo, Lining
Skokut, Thomas A.
Young, Scott A.
Folkerts, Otto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NOMBER: 0.5000, 1996
CLASSIFICATION: 000
RIOR APPLICATION BOOD
RIOR APPLICATION DATA: 0.001,135
APPLICATION NUMBER: 06/001,135
APPLICATION NUMBER: 08/300,726
PILING DATE: 3ceptember 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard,
REGISTRATION NUMBER: 32,32
REFERENCE/DOCKET NUMBER: 219/247
TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: MCSWiggen, James A. APPLICANT: McSwiggen, James A. APPLICANT: McSwiggen, James A. APPLICANT: McSwiggen, James A. APPLICANT: McSwiggen, Patricia Ann Owe APPLICANT: Skokut, Thomas A. APPLICANT: Skokut, Thomas A. APPLICANT: Poung, Scott A. APPLICANT: Poung, Scott A. APPLICANT: Poleste, Otto APPLICANT: Poleste, Otto APPLICANT: McTo, Donald J. TITLE OF INVENTION: MODULATION OF TITLE OF SEQUENCES: 1263
CORRESPONDENCE ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                 RESULT 382
US-08-679-645-147
; Sequence 147, Application US/08679645
; Patent No. 6350934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 147:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                     941 AGAGGTGTGAGCGCAGA 957
                                                                                                                                                                                                                                                                                1 AGAGGUAUCAGAGCAGA 17
                      LENGTH: 17 base pairs;
TYPE: nucleic acid
STRANDEDNESS: single
// TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
               SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Zwick,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;
TOPOLOGY:
US-08-679-645-147
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Gaps

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Query Match 0.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 23.5%; Pred. No. 2.1e+02; Matches 4; Conservative 10; Mismatches 3; Indels
                                                                                                                                                                                                                                                           APPLICANT: GARAGES, DOMAIG J.

APPLICANT: MARTON: COMPOSITION AND METHODS FOR TITLE OF INVENTION: MODULATION OF GENE EXPRESSION TITLE OF INVENTION: MODULATION OF GENE EXPRESSION TITLE OF INVENTION: IN PLANTS
NUMBER OF SEQUENCES: 1263
CORRESSERE: Lyon & Lyon
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZITY: Los Angeles
STATE: Galifornia
COUNTRY: U.S.A.
ZITY: P9071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: GOMPATA:
MEDIUM TYPE: J.S.A.
ZITY: MEDIUM TYPE: J.S.A.
SOFTWARE: MOOT Perfect 5.1
COMPUTER: IBM Compatible
OPERATING SYSTEM: BNO TO PERfect 5.1
COMPUTER: JULY 12, 1996
CLASSIFICATION NUMBER: US/08/679,645
FILING DATE: JULY 13, 1995
RILING DATE: JULY 13, 1995
RILING DATE: SEQUENDER 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: WARNATION:
NAME: WARNATION TO AND TO AND TO THE T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REPERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELEPRONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                Edington, Brent E.
McSwiggen, James A.
Merlo, Patricia Ann Owens
Guo, Lining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-474-432B-355/c
; Sequence 355, Application US/09474432B
; Patent No. 6528640
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1105 TATGTAGTTTTCTGTTT 1121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : :|:| ::::|:|
1 UUUGUAUUUUUCUGCUU 17
                                                                                                                                                                     Skokut, Thomas A.
Young, Scott A.
Folkerts, Otto
Merlo, Donald J.
Zwick, Michael G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-679-645-888
                                                                       APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 386
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0.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 23.5%; Pred. No. 2.1e+02;
Matches 4; Conservative 10; Mismatches 3; Indels
                                                                                                                      Gequence 887, Application US/08679645

Sequence 887, Application US/08679645

Patent No. 635034

GENERAL INFORMATION:
APPLICANT: Edington, Brent E.
APPLICANT: McSwiggen, James A.
APPLICANT: McSwiggen, James A.
APPLICANT: McSwiggen, James A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Folkerts, Otto
APPLICANT: McTlo, Donald J.
TITLE OF INVENTION: McDULATION OF GENE EXPRESSION
TITLE OF INVENTION: MCDULATION OF GENE EXPRESSION
TITLE OF INVENTION: IN PLANTS
NUMBER OF SEQUENCES: 1263

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
CLASSIFICATION: 800
PRIOR APPLICATION B00
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/001,135
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER RADABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ALICHAEL MACHALLY
NAME: WALDLEY, RICHAELD
REGISTRATION NUMBER: 32,327
REPERRENCE/DOCKET NUMBER: 219/2
REPERRENCE/DOCKET NUMBER: 219/2
RELECHONE: (213) 489-1600
TELEPHONE: (213) 955-0440
TELERA: (213) 955-0440
TELERA: (7-3510
INFORMATION FOR SEQ ID NO: 887: SEQUENCE CHARACTERISTICS: LENGTH: 17 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1104 TTATGTAGTTTTCTGTT 1120
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US-08-679-645-887
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Gaps

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GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Beigelman, Leo
APPLICANT: Burgin, Alex
APPLICANT: Beaudry, Amber
APPLICANT: Karpelsky, Alex
APPLICANT: Karpelsky, Alex
APPLICANT: Sweedler, David

RESULT 385
US-08-679-645-888
; Sequence 888, Application US/08679645
; Patent No. 6350934
; GENERAL INFORMATION:

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US-09-474-432B-005
Sequence 605, Application US/09474432B
Sequence 605, Application US/09474432B
Sequence 605, Application US/09474432B
Sequence 605, Application US/09474432B
GENERAL INFORMATION:
APPLICANT: Beigelman, Leo
APPLICANT: Beaudry, Amber
APPLICANT: Beaudry, Amber
APPLICANT: Arrelsky, Amber
APPLICANT: Stanen, Shaw
APPLICANT: Stanen, Shaw
APPLICANT: Zinnen, Shaw
APPLICANT: Zinnen, Shaw
APPLICANT: Zinnen, Shaw
APPLICANT: Shaw
FILING DATE: 1999-12-19
PRIOR FILING DATE: 1999-12-19
PRIOR FILING DATE: 1998-11-04
PRIOR FILING DATE: 1998-04-29
PRIOR FILING DATE: 1998-04-29
PRIOR FILING DATE: 1999-04-28
PRIOR FILING DATE: 1999-04-28
NUMBER OF SEQ ID NOS: 1226
SEQ ID NO 605
LENGTH. 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Beigelman, Leo
APPLICANT: Beigelman, Leo
APPLICANT: Beigelman, Leo
APPLICANT: Berdard, Amber
APPLICANT: Berdard, Amber
APPLICANT: Bardary, Amber
APPLICANT: Karpeisky, Alex
APPLICANT: Adamic, Jasenka
APPLICANT: Sweedler, David
APPLICANT: Sinnen, Slawn
TITLE OF INVENTION: Nucleotide triphosphate and their incorporation into oligonuclec
FILE OF INVENTION UNMERR: US 60/064,866
PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR APPLICANTION NUMBER: US 60/064,727
PRIOR FILING DATE: 1999-04-28
PRIOR FILING DATE: 1999-04-28
PRIOR FILING DATE: 1999-04-28
NUMBER OF SEQ ID NOS: 1526
SOOTWARE: Patentin version 3.0
SEQ ID NO 727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 727, Application US/09474432B
Patent No. 6528640
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              523 CTGCCGGAGGAGCAGCT 539
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Best Local Similarity 70.67
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Homo sapiens
US-09-474-432B-727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
CRGANISM: Homo sapiens
US-09-474-432B-605
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APPLICANT: Bandiry, Amber
APPLICANT: Bandiry, Alex
APPLICANT: Karpeisky, Alex
APPLICANT: Samedier, David
APPLICANT: Sweedler, David
APPLICANT: Sweedler, David
APPLICANT: Sweedler, David
APPLICANT: Zinnen, Shawn
ITITE OF INVENTION NUMBER: US/09/474,432B
CURRENT FILING DATE: 1999-12-19
CURRENT APPLICATION NUMBER: US 60/064,866
PRIOR FILING DATE: 1999-12-19
PRIOR FILING DATE: 1999-12-19
PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR PLING DATE: 1999-04-29
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR FILING DATE: 1999-04-28
NUMBER: US 09/186,675
PRIOR FILING DATE: 1999-04-28
NUMBER: PREDERICATION NUMBER: US 09/301,511
PRIOR FILING DATE: 1999-04-28
NUMBER OF SEQ ID NOS: 1526
SOFTWARE: PATCHTIN VERSION 3.0
APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleotide triphosphate and their incorporation into oligonucleot
FILE REFERENCE: MelHolo-83-18 (247/276)
CURRENT APPLICATION NUMBER: US/09/474,432B
CURRENT FILING DATE: 1999-12-19
PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR APPLICATION NUMBER: US 60/064,727
PRIOR FILING DATE: 1998-04-29
PRIOR FILING DATE: 1998-04-29
PRIOR FILING DATE: 1998-11-04
PRIOR FILING DATE: 1999-04-28
NUMBER OF SEQ ID NOS: 1526
SOFTWARE: Patentin version 3.0
SEQ ID NO 355
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 2.1e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 479, Application US/09474432B
; Patent No. 6528640
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1240 CTGGACGTGGCCATGTG 1256
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Best Local Similarity 58.8
Matches 10; Conservative
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Burgin, Alex
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US-09-474-432B-479
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: RNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 387
US-09-474-432B-479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-474-432B-355
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    à
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RESULT 388

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                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 104, Application US/09265503B
Fatent No. 6538108
GENERAL INFORMATION:
APPLICANT: Liskay, Robert M.
APPLICANT: Baconner, C. Eric
APPLICANT: Baker, Sean M.
APPLICANT: Ballag, Ronn J.
APPLICANT: Rolodner, Richard D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: RELATING TO DNA MISMATCH REPAIR GENES
NUMBER OF SEQUENCES: 148
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kolisch, Hartwell, Dickinson, McCormack & Heuser
                                                                                                                                                                                                                                                                                                   Query Match

0.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 76.5%; Pred. No. 2.1e+02;
Matches 13; Conservative 1; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PSYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.9%; Score 12.2; DB 1; 82.4%; Pred. No. 2.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Van Rysselberghe, Pierre C.
REGISTRATION NUMBER: 33,557
REFERENCE/DOCKET NUMBER: OHSU 306D
TELECOMMUNICATION INFORMATION:
TELEPHONE: (503) 224-6655
TELEPHONE: (503) 295-6679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/265,503B
FILING DATE: March 10, 1999
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR FILING DATE: 1998-11-04
PRIOR APPLICATION NUMBER: US 09/301,511
PRIOR PILING DATE: 1999-04-28
NUMBER OF SEQ ID NOS: 1526
SOFTWARE: PatentIn version 3.0
SEQ ID NO 831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (503) 295-6679
TELEX: 360619
INFORMATION FOR SEQ ID NO: 104:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                         894 GCGGTACAGCGTGGCCC 910
                                                                                                                                                                                                                                                                                                                                                                                                                                           GCGGUACAGUGAGGACC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                        TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Portland
STATE: Oregon
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-265-503B-104/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-265-503B-104
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Beaudiry, Amber
APPLICANT: Beaudiry, Amber
APPLICANT: Beaudiry, Amber
APPLICANT: Karpeisky, Alex
APPLICANT: Karpeisky, Alex
APPLICANT: Adamic. Jasenka
APPLICANT: Sweedler, David
APPLICANT: Sweedler, David
APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleoctide triphosphate and their incorporation into oligonucleoct
TITLE OF INVENTION: Nucleoctide triphosphate and their incorporation into oligonucleoct
FILE REFERENCE: MBH800-831-B (247/276)
CURRENT APPLICATION NUMBER: US 60/064,866
PRIOR PLING DATE: 1999-11-05
PRIOR PLING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR PLING DATE: 1998-04-29
PRIOR PLING DATE: 1999-04-28
NUMBER: US 09/186,675
PRIOR FILING DATE: 1999-04-28
NUMBER: OF SEQ ID NOS: 1526
SOFTWARE: PARCENTIN VUMBER: US 09/301,511
SEQ ID NO 776
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Karpeisky, Alex
APPLICANT: Karpeisky, Alex
APPLICANT: Adamic, Jasenka
APPLICANT: Sweedler, David
APPLICANT: Sannen, Shawn
TITLE OF INVENTION: Nucleotide triphosphate and their incorporation into oligonucleot
FILE REPERBURG: MBHB00-831-B (247/276)
CURRENT APPLICATION UNMER: US/09/474,432B
CURRENT FILING DATE: 1999-12-19
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                               Score 12.2; DB 1; Length 17; Pred. No. 2.1e+02; 4; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR FILING DATE: 1997-11-05
PRIOR APPLICATION NUMBER: US 60/004,727
PRIOR FILING DATE: 1998-04-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 831, Application US/09474432B Patent No. 6528640
                                                                                                                                                                                                                                                                                                        Sequence 776, Application US/09474432B
Patent No. 6528640
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Burgin, Alex
Beaudry, Amber
                       Ouery Match
Best Local Similarity 58.84
Matches 10; Conservative
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Burgin, Alex
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CORGANISM: Homo sapiens
US-09-474-432B-776
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Best Local Similarity
Matches 12; Conserv
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US-09-474-432B-831
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APPLICANT:
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APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Jan
APPLICANT: Stinchcomb, Jan
APPLICANT: Stinchcomb, Jan
APPLICANT: Stoched, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions F
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCES: MHERE: US/09/371,72B
CURRENT FILING DATE: 1999-08-10
PRIOR PILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin Version 3.0
SEQ ID NO 573
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Pred. No. 2.1e+02;
3; Mismatches 3; Indels
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Pred. No. 2.1e+02;
0; Mismatches 3;
                                                                                                                                                                     APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                         Sequence 573, Application US/09371772B Patent No. 6566127 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                843 AGATGGGTCAGCATACC 859
             432 CACGIICAGAAAGIIGC 448
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1 Similarity 64.7%;
11; Conservative
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Best Local Similarity 82.4
Matches 14; Conservative
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US-09-371-772B-769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: RNA
ORGANISM: Homo sapiens
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Best Local Similarity
Matches 11; Conserv
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US-09-371-772B-467/c
; Sequence 467, Application US/09371772B
; Patent No. 6566127
; GENERAL INFORMATION:
; APPLICANT: Ribozyme Pharmaceuticals, Inc.
; APPLICANT: Bavozyme Pharmaceuticals, Inc.
; APPLICANT: Bavozyme Pharmaceuticals, Inc.
; APPLICANT: Stindboomb, Dan
; APPLICANT: Scobedo, Jaime
; APPLICANT: Scobedo, Jaime
; APPLICANT: Scobedo, Jaime
; TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
; TITLE OF INVENTION: Wetels of Vascular Endothelial Growth Factor Receptor
; FILE REFERENCE: MBHBO0, 876-J (237/198)
; CURRENT APPLICATION NUMBER: US/09/371,772B
; PRIOR APPLICATION NUMBER: US 60/005,974
; PRIOR APPLICATION NUMBER: US 60/005,974
; PRIOR FILING DATE: 1996-01-08
; NUMBER OF SEQ ID NOS: 14225
; SOFTWARE PATENTIN VERSION 3.0
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0.9%; Score 12.2; DB 1; Length 17;

Best Local Similarity 58.8%; Pred. No. 2.1e+02;

Matches 10; Conservative 4; Mismatches 3; Indels
   Indels
   Mismatches
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                                    941 AGAGGTGTGAGCGCAGA 957
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                                                                17 AGACGTGAGAGCCCAGA
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Best Local Similarity 82.4
Matches 14; Conservative
     Conservative
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US-09-371-772B-454
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US-09-371-772B-454
     14;
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     Matches
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APPLICANT: ALCANT: Pavoco, Pam
APPLICANT: Pavoco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Scobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (227/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
FRIOR PAPLICATION NUMBER: US 60/005,974
FRIOR PAPLICATION NUMBER: US 08/584,040
FRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 142255
SOFTWARE: Patentin version 3.0
SEQ ID NO 1607
LENGTH: 17

WANTE: DATE

LENGTH: 17
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APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stindcomb, Dan
APPLICANT: Stindcomb, Dan
APPLICANT: Stindcomb, Dan
APPLICANT: Stindcomb, Dan
APPLICANT: Stockedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions F
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHBO, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371, 772B
CURRENT FILING DATE: 1995-08-10
FRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARR: Patentin version 3.0
SEQ ID NO 1678
LENGTH...1
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Similarity 23.5%; Pred. No. 2.1e+02;
4; Conservative 10; Mismatches 3;
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; Sequence 1679, Application US/09371772B
; Patent No. 650, Application US/09371772B
; GENERAL INFORMATION;
; APPLICANT: Ribozyme_Pharmaceuticals, Inc.
                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                       Sequence 1607, Application US/09371772B Patent No. 6566127
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; ORGANISM: Homo sapiens
US-09-371-772B-1607
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Best Local Similarity
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US-09-371-772B-1607
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APPLICANT: Saturchecomb, Jam
APPLICANT: Stinchcomb, Dan
APPLICANT: Bacobedo, Jain
TITLE OF INVENTION: Method and Readent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR PELING DATE: 1995-01-08
PRIOR PELING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEG ID NO 1506
LENGTH: 17
                                                                                                                                                                                                                                                                                                                 APPLICANT: RGSW-19gen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Scobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of V337/198)
CURRENT FILING DATE: 1999-08-10
PRIOR FILING DATE: 1999-08-10
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEQ ID NO 1078
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58.8%; Pred. No. 2.1e+02;
Ive 4; Mismatches 3; Indels
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Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
                                                                                                                                                                                                                              APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                                                                                ; Sequence 1078, Application US/09371772B
; Patent No. 6566127
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Best Local Similarity 58.8
Matches 10; Conservative
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US-09-371-772B-1506
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ORGANISM: Homo sapiens
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US-09-371-772B-1506
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US-09-371-772B-1078
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Sequence 3069, Application US/09371772B

Sequence 3069, Application US/09371772B

Sequence 3069, Application US/09371772B

Sequence 3069, Application US/09371772B

Patent No. 6566127

GENERAL INFORMATION:

APPLICANT: Ravoo, Pam

APPLICANT: Ravoo, Pam

APPLICANT: Barobedon, Jam

APPLICANT: Escobedon, Jam

CURRENT APPLICANTION: Method and Reagent for the Treatment of Diseases or Conditions of TITLE OF INVENTION: Lavels of Vascular Endothelial Growth Factor Receptor

FILE REFERENCE: MBHB00,876-31(237,198)

CURRENT APPLICATION NUMBER: US 60/005,974

PRIOR PILING DATE: 1996-10-26

PRIOR PILING DATE: 1996-10-8

NUMBER OF SEQ ID NOS: 14225

SOFTWARE: PatentIn version 3.0

SEQ ID NO 3069

LENGTH: 17
                                                                                                                                                          APPLICANT: Raveo, Faun
APPLICANT: Struchcomb, Dan
APPLICANTON Wethod and Reagent for the Treatment of Diseases or Conditions F
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
CURRENT APPLICATION NUMBER: 1899-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-01-26
PRIOR PILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn Version 3.0
SSQ ID NOS: 2764
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 2.1e+02;
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82.4%; Pred. No. 2.1e+02;
ive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
Sequence 2764, Application US/09371772B Patent No. 6566127 CENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        878 CCAAGITCCAGGAGCIG 894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    881 AGTICCAGGAGCTGCGG 897
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Best Local Similarity
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US-09-371-772B-3069/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: RNA
; ORGANISM: Mus sp.
US-09-371-772B-2764
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Best Local S
Matches 14
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                         Sequence 1679, Application US/09371772B

Sequence 1679, Application US/09371772B

Sequence 1679, Application US/09371772B

GENERAL INFORMATION:
APPLICANT: Bibozyme Pharmaceuticals, Inc.
APPLICANT: Bibozyme Pharmaceuticals, Inc.
APPLICANT: Bichocyme Pauco, Dam
APPLICANT: Estobedo, Jame
APPLICANT: Estobedo, Jame
APPLICANT: Estobedo, Jame
APPLICANT: Bichocyme Sequence Conditions Re
ITILE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
ITILE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
ITILE REPERBUCE: MBHB00, 876-J (237/198)
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1679
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UG-09-371-772B-2697/C

Sequence 2697, Application US/09371772B

Patent No. 5666127

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: Bacobed, Jam
APPLICANT: Escobed, Jam
APPLICANT: Escobed, Jam
APPLICANT: Brobed, Jam
APPLICANT: Brobed, Jam
APPLICANT: Brobed, Jam
APPLICANT: Brobed, Jam
APPLICANT: McSwigen, Jam
APPLICANT: Brind Darie: 1999-00-10
FRIOR FILING DATE: 1999-01-08
PRIOR FILING DATE: 1995-01-08
PRIOR FILING DATE: 1995-01-08
SRIOR APPLICANTON NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-01-08
SRIOR WARREN FILING DATE: 1995-01-08
SRIOR FILING DATE: 1995-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 12.2; DB 1; Length 17; 29.4%; Pred. No. 2.1e+02; ive 9; Mismatches 3; Indels
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Best Local Similarity 29.4
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TYPE: RNA
; ORGANISM: Mus sp.
US-09-371-772B-2697
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                         US-09-371-772B-1679
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GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Bavco, Pam
APPLICANT: Stinchcomb, Dan
APPLICANT: British and Statement of Diseases or Conditions F
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions F
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions F
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: MUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR PPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin Version 3.0
SEQ ID NO 3660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Baccobed, Jam
APPLICANT: Bacchedo, Jame
APPLICANT: Bacchedo, Jame
APPLICANT: Bacchedo, Jame
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MelhBoo, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-36
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEQ ID NO 4161
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4161, Application US/09371772B Patent No. 6566127
                                           Sequence 3660, Application US/09371772B Patent No. 6566127
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Best Local Similarity 70.6'
Matches 12; Conservative
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Best Local Similarity 70.6
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TYPE: RNA
; ORGANISM: Mus sp.
US-09-371-772B-3660
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ORGANISM: Mus sp.
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RESULT 407
US-09-371-772B-3660
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US-09-371-772B-3387
; Sequence 3387, Application US/09371772B
; Patent No. 6566127
; GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavoo, Pam
APPLICANT: Excobedo, Jaime
APPLICANT: Stinchcomb, Dan
APPLICANT: Scobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: NBH800,876-J (2377198)
CURRENT APPLICATION NUMBER: US/09/371,772B
                                                                                                                                                                           APPLICANT: Pavco, Pam
APPLICANT: McWaiggen, Jim
APPLICANT: Stinchcoub, Dan
APPLICANT: Stinchcoub, Dan
APPLICANT: Stinchcoub, Dan
APPLICANT: Stinchcoub, Dan
APPLICANT: Secobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE SEPERANCE: MBHB00, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR PLING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEQ ID NO 3213
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 12.2; DB 1; Length 17; Pred. No. 2.1e+02; 0; Mismatches 3; Indels
                                                                      Sequence 3213, Application US/09371772B
Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Rayco, Pam
APPLICANT: McSwiggen, Jim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3387
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Dest Local Similarity 82.4%;
Matches 14; Conservative
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Best Local Similarity 52.9
Matches 9; Conservative
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; ORGANISM: Mus sp.
US-09-371-772B-3387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ) ORGANISM: Mus sp. US-09-371-772B-3213
                                                   US-09-371-772B-3213/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
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RESULT 409

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USOURTH 1-172B-5116/C

Sequence 5116, Application US/09371772B

Sequence 5116, Application US/09371772B

Patent No. 6566127

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Ravoco, Pam

APPLICANT: Stinchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Scobedo, Jaime

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions I TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions I TITLE OF INVENTION: MABER: US/09/371,772B

CURRENT APPLICATION NUMBER: US 60/005,974

PRIOR PILING DATE: 1995-08-10

PRIOR PILING DATE: 1996-01-08

NUMBER OF SEQ ID NOS: 14225

SEQ ID NOS: 14225

SEQ ID NO 5116
                                                                                         APPLICANT: Pavco, Pam
APPLICANT: Braco, Jam
APPLICANT: Stinchcomb, Jim
APPLICANT: Stinchcomb, Jaim
APPLICANT: Stinchcomb, Jaime
APPLICANT: Stinchcomb, Jaime
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00, 876-J (237/198)
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-06
RIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin Version 3.0
SEQ ID NO 4222
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Conservative 3; Mismatches 3; Indels
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82.4%; Pred. No. 2.1e+02;
tive 0; Mismatches 3;
                       Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
4722, Application US/09371772B
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Best Local Similarity 82.4
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Homo sapiens
US-09-371-772B-4722
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Best Local Similarity
Matches 11; Conserv
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Sequence 4643, Application US/09371772B

Sequence 4643, Application US/09371772B

Sequence 4643, Application US/09371772B

SERVERAL INFORMATION:

APPLICANT: Rabozyme Pharmaceuticals, Inc.

APPLICANT: Baveco, Pau

APPLICANT: Stinchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Stochcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANTION: Levels of Vascular Endothelial Growth Factor Receptor

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

TITLE OF INVENTION NUMBER: US/09/371,772B

CURRENT APPLICATION NUMBER: US 60/005,974

PRIOR FILING DATE: 1995-08-10

PRIOR APPLICATION NUMBER: US 60/005,974

PRIOR FILING DATE: 1995-01-08

NUMBER OF SEQ ID NOS: 14225

SEQ ID NOS: 14225

SEQ ID NO 4643
                                                                                            GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Rosuggen, Jum

APPLICANT: McSwiggen, Jum

APPLICANT: StinchComb, Dan

APPLICANT: StinchComb, Dan

APPLICANT: StinchComb, Dan

APPLICANT: Escobedo, Jaime

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION: Method and Reagent Endothelial Growth Factor Receptor

TITLE OF INVENTION WIMERS: US 09/371,772B

CURRENT APPLICATION NUMBER: US 60/005,974

PRIOR FILING DATE: 1999-08-10-26

PRIOR FILING DATE: 1996-01-08

NUMBER OF SEQ ID NOS: 14225

SOFTWARE: PatentIn version 3.0

LENGTH: 17
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       US-09-371-772B-4457/c
; Sequence 4457, Application US/09371772B
; Patent No. 6566127
; GENERAL INFORMATION:
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Best Local Similarity 82.4
Matches 14; Conservative
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US-09-371-772B-4457
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US-09-371-772B-4643/c
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; Sequence 5579, Application US/09371772B

RESULT 413 US-09-371-772B-5579/c

> RESULT 411 US-09-371-772B-4722

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GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Rabozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Danine
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions F
TITLE OF INVENTION: McMeder: US/09/371,772B
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT APPLICATION NUMBER: US/09/371,772B
FILE REPERENCE: MSHBO0, 976-10
PRIOR APPLICATION NUMBER: US/09/371,772B
PRIOR PELING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
SGOTWARE: Patentin version 3.0
LENGTH: 17
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APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MEBBOO, 876-J (227/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-10-26
PRIOR FILING DATE: 1999-10-26
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 6624
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0.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 2.1e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
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; Sequence 6624, Application US/09371772B
; Partent No. 656613.
; GRNERAL INFORMATION:
; APPLICANT: Ribozyme Pharmaceuticals, Inc.
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's Sequence 6701, Application US/09371772B

; Patent No. 6566127

; GENERAL INFORMATION.
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CRGANISM: Homo sapiens
US-09-371-772B-6624
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APPLICANT: Stinchtoomb, Dan
APPLICANT: Bacobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MEHBOO, 187-0/ (237/198)
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 68/584,040
PRIOR APPLICATION NUMBER: US 68/584,040
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
                                                                                                                                                      APPLICANT: ROSWIGGEN, Jim
APPLICANT: ACRAUGH TO TEACH MICHORAN ACRAUGH TO TEACH MICHORAN ACRAUGH TO TEACH TO TE
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Pred. No. 2.1e+02;
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Best Local Similarity 82.4%; Pred. No. 2.1e
Matches 14; Conservative 0; Mismatches
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
               Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-371-772B-6439/c
; Sequence 6439, Application US/09371772B
; Patent No. 6566127
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; Patent No. 6566127
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Best Local Similarity 29.4%;
Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                   Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1146 TTTTCT
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US-09-371-772B-6296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-371-772B-5579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 6296
LENGTH: 17
                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  원
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Blater, Michael R.
APPLICANT: Huang, Farinett, James R.
APPLICANT: Bolchakova, Elena
APPLICANT: Bolchakova, Elena
APPLICANT: Storts, Douglas R.
APPLICANT: Storts, Douglas R.
APPLICANT: OF INVENTION: THERMOPHILIC DNA POLYMERASES FROM
TITLE OF INVENTION: THERMOPORA NEAPOLITANA
TITLE OF INVENTION: THERMOTOGA NEAPOLITANA
TITLE OF INVENTION: THERMOTOGA NEAPOLITANA
TITLE OF INVENTION: THERMOTOGA NEAPOLITANA
TITLE OF SEQUENCES:
TITLE OF INVENTION: THERMOTOGA NEAPOLITANA
TITLE OF INVENTIONA
THERMOTOGA NEAPOLITANA
THERMOTOGA NEAPOLITANA
THERMOTOGA NEAPOLIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 17
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MEDIUM TYPE: Riopy disk
COMPUTER: INPW PC compatible
COMPUTER: INPM PC compatible
APPLICATION NUMBER: PC/US96/09641
FILING DATE: 31-MAY-1996
CLASSIFICATION:
NAME: INPORMATION:
NAME: INPORTATION:
REGISTRATION NUMBER: 40.027
REPERBNICK/DOCKET NUMBER: PRMG-02185
FELECOMMUNICATION:
TELEPHONE: (415) 705-8410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match

0.9%; Score 12.2; DB 1;
Best Local Similarity 82.4%; Pred. No. 2.1e+02;
Matches 14; Conservative 0; Mismatches 3;
                                                                                             4600-0202.41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ANTI-SENSE: NO
ORIGINAL SOURCE:
INDIVIDUAL ISOLATE: Primer PGEX-R
PCT-US95-06266-87
                         NAME: Fabian, Gary R.
REGISTRATION NUMBER: 33,875
REFERENCE/DOCKET NUMBER: 4600
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0860
INFORMATION FOR SEQ ID NO: 87: SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            625 GACCAGCTCCAGGAGCT 641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 GACCGTCTCCGGGAGCT 17
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                                                                                                                                                                                                                                                                                                                                                                                     LOFOLOGY: linear
MOLECULE TYPE: CDNA to MRNA
HYPOTHETICAL: NO
ANTH-CEDAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (415) 397-8338
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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TOPOLOGY: li
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                                                                    APPLICANT: ROWING Fau Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Jame
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
APPLICANTION INVENT: US/09/371,772B
CURRENT APPLICATION NUMBER: US/09/371,772B
PRIOR APPLICATION NUMBER: US/09/371,772B
PRIOR APPLICATION NUMBER: US/09/371,772B
PRIOR APPLICATION NUMBER: US/09/371,772B
PRIOR APPLICATION NUMBER: US/09/005,974
PRIOR FILING DATE: 1995-10-26
RUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detection of Viral Antigens Coded
by Reverse Reading Frames
157
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/06266
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Detection of Viral Ant TITLE OF INVENTION: by Reverse Reading Fr: NUMBER OF SEQUENCES: 157
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Associates
STREET: 350 Cambridge Avenue, Suite 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/246,985
FILING DATE: 20-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/285,561
FILING DATE: 03-MG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/329,729
FILING DATE: 26-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/344,271
FILING DATE: 13-MC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/357,509
FILING DATE: 16-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/357,509
FILING DATE: 15-FEB-1995
          APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 87, Application PC/rUS9506266 GENERAL INFORMATION:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               260 TCCTGGGCTGGCTGATC 276
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 82.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: US
ZIP: 94306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 418
PCT-US95-06266-87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 6701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: P
STATE:
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                    Length 17;
                                                                3; Indels
                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: StinchComb, Dan T.
APPLICANT: StinchComb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: NETHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                          TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL GROWTH FACTOR
                0.9%; Score 12.2; DB 1;
82.4%; Pred. No. 2.1e+02;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 8.0-rage
COMPUTER: 1BM Compatible
OPERATING SYSTEM: 1BM P.C. DOS 5.0
SOFWARE: WORD PRIFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY AGENT INFORMATION:
NAME: WALDIES, RICHARD 3.327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
NAME: WALDIES, RICHARD 3.327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEBOONE 1.213 4.99-1600
                                                                                                                                                                                                                                         US-08-584-040-3044/c
; Sequence 3044, Application US/08584040
; Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Lyon & Lyon
633 West Fifth Street
                                                                                                            1199 GACCTTCACACCTCCCC 1215
                                                                                                                                               1 daccirdacacaccic 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (213) 489-160
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 82.4<sup>5</sup>
Matches 14; Conservative
Query Match
Best Local Similarity 82...
Thes 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
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1027 CAAAGTGCAGCTGACTC 1043

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18 CAAAAAGCAGCTGGCTC 2

US-09-371-772B-1472/c

RESULT 421

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APPLICANT: McSwiggen, Jam
APPLICANT: Stinchcomb, Dan
APPLICANT: Stochedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions F
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MHBBO0, 876-J (237/198)
CURRENT FILING DATE: 1999-08-10
PRIOR PILING DATE: 1995-08-10
PRIOR PILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTHARE: PatentIn version 3.0
SEQ ID NO 1472
LENGTH: 18
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| Patent No. 5596091
| GENERAL INFORMATION:
| APPLICANT: SWITZER, Christopher
| TITLE OF INVENTION: NOVEL ANTISENSE OLIGONUCLEOTIDES
| NUMBER OF SEQUENCES: 13
| CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: R10py disk
MEDIUM TYPE: R10py disk
COMPUTER: IBM PC compatible
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/214,603
FILING DATE: 18-MAR-1994
CLASSIFICATION: 536
ATTORNEY/AGENTINORMANIANDEM: 3369
REGISTRATION NUMBER: 37.369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Townsend and Townsend Khourie and Crew STREET: Steuart Street Tower, One Market Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Ub ... 2.4e+02; 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
0.9%; Score 12.2; DB 1;
Best Local Similarity 82.4%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 2307E-052100US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
TELEFAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 11:
                      Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme_Pharmaceuticals, Inc.
Sequence 1472, Application US/09371772B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1027 CAAAGTGCAGCTGACTC 1043
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LENGTH: 12 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: Steuart Stre
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                            Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 422
US-08-214-603-11/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                       APPLICANT:
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Pred. No. 1.2e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCT-US95-06379-14/c

PCT-US95-06379-14/c

Sequence 14, Application PC/TUS9506379

GENERAL INFORMATION:

APPLICANT: Watanabe, Kyoichi A.

APPLICANT: Ren, Win-Yun

ADDRESSEE: Cooper & Dunham ILP

ADDRESSEE: Cooper & Dunham ILP
                                                                                                                                                                                                                                                                                                                                      44683-Z/JPW/MJG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.9%; Score 12; DB Best Local Similarity 100.0%; Pred. No. 1.2 Matches 12; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUBER: PCT/US95/06379
FILING DATE: May 13, 1994
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
                                                                                                                                            COMPUTER: 1BM PC
COMPUTER: 1BM PC
COMPUTER: 1BM PC
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,138
FILING DATE: Unne 7, 1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 44683-Z/JPW
TELECHOWNICATION INFORMATION:
TELECHORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGHH. 13 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2: Cooper & Dunham LLP
1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 44683-PCT
FELECOMUNICATION INFORMATION:
TELEPHONE: 212-278-0400
TELEPHONE: 212-391-0526
1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch 1.44Mb
COMPUTER: IBM PC
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                       ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch 1.44Mb
AMMITTER: IBM PC PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , MOLECULE TYPE: DNA (genomic)
US-08-484-138-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELBFAX: 212-391-0526
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13 rritriciiri 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 1185 Aven
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                                        New York
                                         New York
                                                                        COUNTRY: U
     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 425
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                                            Score 12; DB 1; Length 12;
Pred. No. 1e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Watanabe, Kyoichi A.
APPLICANT: Watanabe, Kyoichi A.
APPLICANT: Weil, Welvi
APPLICANT: Weil, Roger
TITLE OF INVENTION: Complementary DNA and Toxins
NUMBER OF SEQUENCES: 43
NUMBER OF SEQUENCES: 43
NUMBER OF SEQUENCES: 6
STREEF: 30 Rockefeller Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%; Score 12; DB 1; L4
100.0%; Pred. No. 1.2e+02;
cive 0; Mismatches 0;
                                            Query Match

0.9%; Score 12; DB
Best Local Similarity 100.0%; Pred. No. 1e+
Matches 12; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 14, Application US/08242664
; Patent No. 5571937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1145 TITITICITITI 1156
                                                                                                                         1144 TTTTTTTTTT 1155
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Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: New York
COUNTRY: U.S.A.
ZIP: 10112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                     RESULT 423
US-08-242-664-14/c
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                 US-08-214-603-11
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Gaps

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RESULT 427
US-08-683-839B-15/c
US-08-683-839B-15/c

j Sequence 15, Application US/08683839B

j Patent No. 5744326

j GENERAL INFORMATION:
   APPLICANT: 111, Charles . R. et al.
   TITLE OF INVENTION: Regulatory Sequences To Increase Expression of
   TITLE OF INVENTION: Intronless Genes Containing Near-Consensus Splice Sites
   TUTLE OF INVENTION: Intronless Genes Containing Near-Consensus Splice Sites
   TORRESPONDENCE ADDRESS:
   ADDRESSEE: LAHIVE & COCKFIELD
   STREET: 60 State Street, suite 510
   CITY: Boston
   STATE: Massachusetts
   COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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APPLICANT: MORIYA, Mika
APPLICANT: MATSUI, Hitcshi
APPLICANT: WOKOZEKI, Kenzo
APPLICANT: HIRANO, Seiko
APPLICANT: HIRANO, Seiko
APPLICANT: HIRANO, Seiko
APPLICANT: TZUI, Masakaz
APPLICANT: SUGIMOTO, Masakazu
TITLE OF INVENTION: METHOD OF AMPLIFYING GENE USING
TITLE OF INVENTION: ARTIFICIAL TRANSPOSON
NUMBER OF SEGUENCES: 32
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: DatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/08/683,839B
FILING DATE: 11-MARCH-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.9%; Score 12; DB 1; Le
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Remillard, Jane E.
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: TTI-138
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 428
US-08-674-168-10
; Sequence 10, Application US/08674168
; Patent No. 5804414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 15
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1125 TGAAAAGAGAG 1136
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Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 14 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12 TGAAAAAGAGAG 1
2 GGACTGACCCCT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-683-839B-15
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| Patent No. 5591577
| GENERAL INFORMATION:
| APPLICANT: TRUCHIYA, MAKOTO
| APPLICANT: MINA, KIYOSHI
| TITLE OF INVENTION: MOBILE GENETIC ELEMENT ORIGINATED FROM
| TITLE OF INVENTION: BREVIBACTERIUM STRAIN
| NUMBER OF SEQUENCES: 9
| CORRESPONDENCE BADRESS: 3
| ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT
| STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, FOURTH FLOOR
| CITY: ARLINGTON | STRAIN | STRAIN | STRAIN | STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, FOURTH FLOOR | STREET: WINDOWN | STRAIN | STREET: WINDOWN | STRAIN | STREET: WINDOWN | STRAIN | ST
                                                                                                                                                                                                                                                                                                                                                    0;
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0.9%; Score 12; DB 1; Length 14;
Best Local Similarity 100.0%; Pred. No. 1.5e+02;
Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                             Length 13;
                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/146,010A
FILING DATE: 12-NOV-1993
CLASSIFICATION NUMBER: US/08/146,010A
FILING DATE: 11-NAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: OBLON, NORMAR: 24,618
REFERENCE/DOCKET NUMBER: 24,618
REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10-649-0
TELEPHONE: (703) 413-220
TELEPHONE: (703) 413-220
TELEPAX: (703) 413-220
TELEPAX: (703) 413-220
TELEPAX: (703) 413-220
TELEPAX: 1000 B:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 base pairs
TYPE: nucleic acid
STRANDENESS: double
TYPE: DUCICIS DINA (genomic)
CORTANTSM: Rrevibacterium lactofermentum
                                                                                                                                                                                                                                                                             Query Match
0.9%; Score 12; DB 1; LeBest Local Similarity 100.0%; Pred. No. 1.2e+02; Matches 12; Conservative 0; Mismatches 0;
                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                           1145 TTTTTCTTTT 1156
              SEQUENCE CHARACTERISTICS:
                                              LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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                                                                                                                                             linear
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US-08-146-010A-8
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US-08-146-010A-8
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963 GGACTGACCCCT 974

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ZIP: 53202-4497
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURREMY APPLICATION DATA:
APPLICATION NUMBER: US/08/365,189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/132,648
FILING DATE: October 5, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Ryser, David G.
REGISTRATION NUMBER: 36,407
REFERENCE/DOCKET NUMBER: 66-005-9367-4
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08.365-189-10

Sequence 10, Application US/08365189
Patent No. 5514576

GENERAL INFORMATION:
APPLICANT: Bower, Patricia A.
TITLE OF INVENTION:
TUTLE OF INVENTION:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Quarles & Brady
STREET: 411 East Wisconsin Avenue
CITY: Milwaukee
    FILING DATE: April 25, 1997
CLASSIFICATION: 800
ATTORNEY/ACENT INFORMATION:
NAME: Gravelle, Micheline
REGISTRATION NUMBER: 40,261
REFERENCE/DOCKET NUMBER: 9369-039
TELECOMMUNICATION INFORMATION:
TELEFAX: (416) 364-7311
TELEFAX: (416) 361-1398
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (414) 277-5717
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (414) 277-571
TELEPAX: (414) 271-3552
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 393 GGCAGCAATGGC 404
                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 100.
Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                   single
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                ; TOPOLOGY; linear;
; MOLECULE TYPE: cDNA
US-08-846-021A-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: cDNA FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                         TYPE: nucleic STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; NAME/KEY:
; LOCATION:
US-08-365-189-10
                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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Fatent No. 5948682
GENERAL INFORMATION:
TITLE OF INVENTION: Preparation of Heterologous Proteins on TITLE OF INVENTION: 011 Bodies
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: BERESKIN & PARR
STREET: 40 King Street West
CITY: Toronto
STREE: Ontario
COUNTRY: Canada
ZIP: MSH 3Y2
COMPUTER: ISM PC compatible
COMPUTER: ISM PC compatible
COMPUTER: ISM PC compatible
COMPUTER: DatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/646,021A
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, ADDRESSEE: P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                       STATE: V. LAGILALA
COUNTRY: USA
ZIP: 2202
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CORRENT APPLICATION DATA:
SOFTWARE: PATENTIN Release #1.0, Version #1.30
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/674,168
FLILNG DATE: 01-UUL-1996
CLASSIFICATION NUMBER: US/08/674,168
FLILNG DATE: 30-UN-1995
APPLICATION NUMBER: US/08/674,168
FLILNG DATE: 30-UN-1996
APPLICATION NUMBER: 24,618
REGISTRATION NUMBER: 24,618
REGISTRATION NUMBER: 10-810-0
TELEFRAN: (703) 413-2220
TELEFRAN: (703) 413-2220
TELEFRAN: (703) 413-220
TELEFRAN: 4703 A13-220
TELEFRAN: 48 DASE DID NO: SEQUENCE CHARACTERISTICS: LENGTH: 14 DASE PAIRS
INPORMATION FOR SED ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 14 DASE PAIRS
ITYPE: mucleic acid
                                         1755 JEFFERSON DAVIS HIGHWAY, SUITE # 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%; Score 12; DB 1; L6
100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Brevibacterium lactofermentum
STRAIN: AJ12036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 100.
Matches 12, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           963 GGACTGACCCCT 974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 GGACTGACCCCT 13
                                                             ARLINGTON : VIRGINIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ANTI-SENSE: NO
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-846-021A-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; STRAIN: A
US-08-674-168-10
                                                                                  STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Dalie, Barbara
APPLICANT: Le, Hung
APPLICANT: Le, Hung
APPLICANT: Miller, Kenneth
APPLICANT: Mirgolo, Nicholas
APPLICANT: Nguyen, Hanh
APPLICANT: Tindall, Stephen
APPLICANT: Zavohny, Paul
TITLE OF INVENTION: Cloning and Expression of
TITLE OF INVENTION: Humanized Monoclonal Antibodies
TITLE OF INVENTION: Against Human Interleukin-4
NUMBER OF SEQUENCES: 69
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schering-Plough Corporation
                                 CORRESPENDENCE ADDRESS:
ADDRESSE: Schering-Plough Corporation
STREET: 2000 Galloping Hill Road
CITY: Kenilworth
STATE: New Jersey
COUNTRY: USA
                                                                                                                                                             COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh
SOFTWARE: Microsoft Word 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/704,744
FILING DATE: 06-SEP-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: ECT/US/95/02400
FILING DATE: 10-MAR-1994
APPLICATION NUMBER: PCT/US/95/02400
FILING DATE: 10-MAR-1994
APPLICATION NUMBER: PCT/US/95/02400
ATTORNEY AGENT INFORMATION:
NAME: FOULK, CYNTHIA L.
REGISTRATION NUMBER: 32,364
REFERENCE/DOCKET NUMBER: 32,364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSE: Schering-Plough Corporation
STREET: 2000 Galloping Hill Road
CITY: Kenilworth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 29, Application US/08469557
Patent No. 5770403
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIF: 07033-0530
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (908) 298-5388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  463 AGCAGCCTGCAG 474
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15 AGCAGCCTGCAG 4
              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 07033-0530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-469-557-29/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-704-744-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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1 Sequence 29, Application US/08208886C

2 Sequence 29, Application US/08208886C

3 Sequence 29, Application US/08208886C

3 Patent No. 559710

3 APPLICANT: Dalle, Barbara

APPLICANT: Murgolo, Nicholas

3 APPLICANT: Tindall, Stephen

5 TILE OF INVENTION: Humanized Monoclonal Antibodies Against Human Interleukin-4

NUMBER OF SEQUENCES: 88

5 CORRESPONDENCE ADDRESS:

5 ADDRESSEE: Schering-Plough Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ..
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                                                      Gaps
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       Score 12; DB 1; Length 15; Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%; Score 12; DB 1; Length 15;
100.0%; Pred. No. 1.8e+02;
Ive 0; Mismatches 0; Indels
                                                  0; Indels
              100.0%; Preq. w.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSE: Schering-Plough Corporation
STREET: 2000 Galloping Hill Road
CITY: Kenilworth
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh
OPERATING SYSTEM: Macintosh
SOFTWARE: Microsoft Word 5.1a
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION TARE: US/08/208,886C
FILING DATE: MARCH 10, 1994
CLASSIFICATION DATA:
APPLICATION UNMBER: TSILING DATE:
APPLICATION NUMBER:
APPLICATION NUMBER:
APPLICATION NUMBER:
APPLICATION NUMBER:
APPLICATION NUMBER:
APPLICATION NUMBER: 32,364
REGISTRATION NUMBER: 32,364
REGISTRATION NUMBER: 32,364
REGISTRATION NUMBER: JB0429
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.5.,
100.0%; Pre
       0.9%;
                                                                                          831 GCTGAAGCTTTC 842
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     463 AGCAGCCTGCAG 474
                                               12; Conservative
                                                                                                                       4 GCTGAAGCTTTC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  in ENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
in TOPOLOGY: linear
US-08-208-886C-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 908 298 5388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15 AGCAGCCTGCAG 4
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 07033-0530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 432
US-08-704-744-29/c
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                                             Matches
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Sequence 62, Application US/08606505B
Setent No. 6114601
Seneral Information:
APPLICANT: KIYOKAWA, Shigeto
APPLICANT: KIYOKAWA, Shigeto
APPLICANT: SHIMADA, Yukihisa
APPLICANT: SHIMADA, Ritsuko
APPLICANT: SHIMADA, Ritsuko
APPLICANT: OKINAKA, Yasushi
TITLE OF INVENTION: NOVEL PLANT GENES
NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
ADDRESSEB: FITZPATRICK, CELLA, HARPER & SCINTO
STREET: 30 ROCKEfeller Plaza
CITY: New YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Le
. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.9%; Score 12; DB Best Local Similarity 100.0%; Pred. No. 1.6 Matches 12; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:

APPLICATION WORBER: US/08/290,793B
FILING DATE: US/08/290,793B
FILING DATE: US/08/290,793B
FILING DATE: 19-FEB-192
APPLICATION NUMBER: US 07/841,659
FILING DATE: 19-FEB-192
APPLICATION NUMBER: US 07/782,784
FILING DATE: 21-MAY-1990
APPLICATION NUMBER: US 07/782,784
FILING DATE: 21-MAY-1990
APPLICATION NUMBER: US 07/499,327
FILING DATE: 21-MAY-1990
APPLICATION NUMBER: US 07/655,966
FILING DATE: 14-FEB-191
APPLICATION NUMBER: US 06/881,553
FILING DATE: 26-OCT-1986
APPLICATION NUMBER: US 06/881,553
FILING DATE: 25-COT-1986
APPLICATION NUMBER: US 06/881,553
FILING DATE: 25-OCT-1986
APPLICATION NUMBER: US 06/881,553
FILING DATE: 25-OCT-1986
APPLICATION NUMBER: US 06/843,958
FILING DATE: 29-NOY-1985
APPLICATION NUMBER: US 06/799,668
FILING DATE: 29-NOY-1985
APPLICATION NUMBER: US 06/799,668
FILING DATE: 25-MAR-1986
APPLICATION NUMBER: US 06/799,668
APPLICATION NUMBER: US 06
                                                                                                                                                                 COMPUTER: Apple Macintosh OPERATING SYSTEM: Macintosh 6.0.5 SOFTWARE: Microsoft Word 5.1A CURRENT APPLICATION DATA:
                                                                     ZIP: 07033-0530
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       463 AGCAGCCTGCAG 474
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15 AGCAGCCTGCAG 4
                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
US-08-290-793B-29
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US-08-606-505B-62
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APPLICANT: Le, Hung
APPLICANT: Miller, Kenneth
APPLICANT: Mirgolo, Nicholas
APPLICANT: Nguyen, Hanh
APPLICANT: Tindall, Stephen
APPLICANT: Zavodny, Paul
TITLE OF INVENTION: Cloning and Expression of
TITLE OF INVENTION: Humanized Monoclonal Antibodies
TITLE OF INVENTION: Against Human Interleukin-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3: Schering-Plough Corporation 2000 Galloping Hill Road
                                                                                                                                          CLASSIFICATION: 424

PRIOR APPLICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: 08/290,793

FILING DATE: ADJUST 16, 1994

APPLICATION NUMBER: PCT/US93/01301

FILING DATE: 19-FEB-1992

APPLICATION NUMBER: US 07/81,659

FILING DATE: 19-FEB-1992

APPLICATION NUMBER: US 07/82,784

FILING DATE: 24-0CT-1991

APPLICATION NUMBER: US 07/499,327

FILING DATE: 21-MAY-1990

APPLICATION NUMBER: US 07/655,966

FILING DATE: 21-0CT-1988

APPLICATION NUMBER: US 07/655,966

FILING DATE: 25-0CT-1987

APPLICATION NUMBER: US 06/881,553

FILING DATE: 25-MAY-1986

APPLICATION NUMBER: US 06/81,553

FILING DATE: 25-MAY-1986

APPLICATION NUMBER: US 06/81,553

FILING DATE: 19-NOV-1986

APPLICATION NUMBER: US 06/81,553

FILING DATE: 19-NOV-1986

APPLICATION NUMBER: US 06/81,553

FILING DATE: 19-NOV-1986
OPERATING SYSTEM: Macintosh 6.0.5
SOFTWARE: Microsoft Word 5.1A
                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,557
FLIANG DATE: 06-UIN-1995
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              32,364
RER: 2409K7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 908-298-5388
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANBEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

0.9%; Soc
Best Local Similarity 100.0%; P.
Matches 12; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
TELEPHONE: 908 298-2987
TELEFAX: 908-298-5388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Foulke, Cynthia L. REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           463 AGCAGCCTGCAG 474
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CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kenilworth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 5863537
GENERAL INFORMATION:
APPLICANT: Dalie,
APPLICANT: Le, Hur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 434
US-08-290-793B-29/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
STREET: 200
CITY: Kenil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-469-557-29
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APPLICANT: SINNETT, Daniel
APPLICANT: SINNETT, Daniel
APPLICANT: LABUDA, Damian
TITLE OF INVENTION: DEFECTION OF CYPIAL, CYP3A4, CYP2D6 AND
TITLE OF INVENTION: MAT2 VARIANTS BY PCR-ALLELE-SPECIFIC OLIGONUCLEOTIDE (ASO)
TITLE OF INVENTION: ASSAY
TITLE OF INVENTION: ASSAY
CURRENT APPLICATION NUMBER: US(90/117,359B
CURRENT FILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 37
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 26
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: New York
STATE: New York
COMPUTRY: U.S.A.
ZIP: 10112-3801
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.50 inch, 720 Kb storage.
COMPUTER: IBM PS/V
OPERATING SYSTEM: MS-DOS Ver3.30
SOFTWARE: PATENT AID Ver1.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/616,990
FILING DATE: 14-311-2000
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: FITZPATRICK, CELLA, HARPER & SCINTO STREET: 30 Rockefeller Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Le
1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KIYOKAWA, Shigeto
SHIMADA, Yukihisa
OHBAXASHI, Masaya
SHIMADA, Ritsuko
OKINAKA, Yasushi
TITLE OF INVENTION: NOVEL PLANT GENES
NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match

0.9%; Score 12; DB
Best Local Similarity 100.0%; Pred. No. 1.6
Matches 12; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: CDNA for use as probes US-09-177-359-26
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FILING DATE: 02-MAR-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Perry, Lawrence S. REGISTRATION NUMBER: 31865 TELECOMMUNICATION INFORMATION:
                                                                                                                       Sequence 26, Application US/09177359B
Patent No. 6183963
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 62, Application US/09616990
Patent No. 6232109
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: KIKUCHI, Yasuhiro
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INFORMATION FOR SEQ ID NO: 62:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    257 ACCTCCTGGGCT 268
  1 TITITICITITI 12
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                                                                                                JS-09-177-359-26/c
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APPLICANT: Chandy, George K.
APPLICANT: Chandy, George K.
APPLICANT: Gurman, George
APPLICANT: Fantino, Emmanuelle
APPLICANT: Fantino, Emmanuelle
APPLICANT: Fantino, Emmanuelle
TITLE OF INVENTION: MARKER AND THERAPEUTIC TARGET
CURRENT APPLICATION NUMBER: US/09/115,446
CURRENT FILING DATE: 1998-07-14
EARLIER APPLICATION NUMBER: 60/052,556
EARLIER FILING DATE: 1997-07-15
BARLIER FILING DATE: 1998-01-08
NUMBER OF SEQ ID NOS: 15
SOFTWARE: FASTSEQ FOR Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%; Score 12; DB 1; Length 15; 100.0%; Pred. No. 1.8e+02;
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                                                                                        720 Kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.9%; Score 12; DB 1; Le Best Local Similarity 100.0%; Pred. No. 1.8e+02; Matches 12; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.9%; Score 12; DB Best Local Similarity 100.0%; Pred. No. 1.8 Matches 12; Conservative 0; Mismatches
STALE: New .....
CUUNTRY: U.S.A.
ZIP: 10112-3801
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.50 inch, 720
COMPUTER: IBM PS/V
COMPUTER: IBM PS/V
COMPUTER: DATENT ALD Verl.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/606,505B
FILING DATE: 23-MRA-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U54963/92
FILING DATE: 02-MRA-1992
ATTORNEY AGENT INFORMATION:
NAME: PERTY, LAWTENCE S.
REGISTRATION NUMBER: 31865
FELCOMMUNICATION INFORMATION:
TELEPHONE: 212-218-2100
JELEPHONE: 212-218-2100
JELEPHONE: 212-218-2100
JELEPHONE: CARRACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: Other nucleic acid
DESCRIPTION: Synthetic DNA
US-08-606-5058-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 3, Application US/09115446
; Patent No. 6165719
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-115-446-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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US-09-115-446-3
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SEQUENCE CHARACTERISTICS:

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GENERAL INFORMATION:
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                                                  US-08-812-951B-2/c
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US-08-812-951B-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 441
US-08-784-747-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Drmanac, Radoje T.
APPLICANT: Drmanac, Snezana
APPLICANT: Drmanac, Snezana
APPLICANT: Hou, Aaron
APPLICANT: Houser, Brian
TITLE OF INVENTION: Detection or Quantification of Nucleic Acid Species
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
STREET: Three Embarcadero Center
STREET: Three Embarcadero Center
CITY: San Francisco
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Pred. No. 1.8e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                   0; Indels
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APPLICATION NUMBER: US/08/812,951B FLING DATE: 04-MAR-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: US/08/784747
FILING DATE: 16-JAN-1997
ATTORNEY/ABATI INFORMATION: NAME: Kumamoto, Andrew A REGISTRATION NUMBER: 40,690
REFERENCE/DOCKTAIN NUMBER: 20411-701
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                        Query Match
0.9%; Score 12; DB
Best Local Similarity 100.0%; Pred. No. 1.8
Matches 12; Conservative 0; Mismatches
                                                         TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid
DESCRIPTION: Synthetic DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 62
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/08812951B Patent No. 6297006
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%;
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Best Local Similarity 92.3
Matches 12; Conservative
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            415-393-2286
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STRANDEDNESS: single
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US-08-812-951B-1
                                                                                                                                    US-09-616-990-62
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STATE:
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                                                                 APPLICANT: Drmanac, Radoje T.
APPLICANT: Drmanac, Snezana
APPLICANT: Brunac, Sanzana
APPLICANT: Houser, Brian
ANTHER OF INVENTION: Detection or Quantification of Nucleic Acid Species
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: MCCUtchen, Doyle, Brown & Enersen LLP
STREET: Three Embarcadero Center
CITY: San Francisco
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Fatent No. 6309824
GENERAL INFORMATION:
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: Methods and Compositions for
TITLE OF INVENTION: Detection or Quantification of Nucleic Acid Species
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: McCutchen, Doyle, Brown & Enersen LLP
STREET: Three Embarcadero Center
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Pred. No. 1.8e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PESESOE FOR Windows Version 2.0 SOFTWARE: PESESOE FOR Windows Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/812,951B
FILING DATE: 04-MAR-1997
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Kumamoto, Andrew A
REGISTRATION NUMBER: 40,690
REFERENCE/DOCKET NUMBER: 20411-701
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US08/784747
FILING DATE: 16-JAN-1997
ATTORNEY AGENT INFORMATION:
; Sequence 2, Application US/08812951B ; Patent No. 6297006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            92.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1146 TTTTTCTTTTGG 1158
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SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 92.3:
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 415-393-20
TELEFAX: 415-393-2286
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TOPOLOGY:
US-08-784-747-3
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/ Sequence 3, Application US/08784747

/ Sequence 3, Application US/08784747

/ GENERAL INFORMATION:

/ APPLICANT: Drymanac, Radoje T.

/ TITLE OF INVENTION: Methods and Compositions for TITLE OF INVENTION: Detection or Quantification of Nucleic Acid Species NUMBER OF SEQUENCES: 13

/ CORRESPONDENCE ADDRESS:

/ ADDRESSEE: McCutchen, Doyle, Brown & Bnersen LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%; Score 12; DB 1; Length 15; 92.3%; Pred. No. 1.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: McCutchen, Doyle, Brown & Enersen LLP
STREET: Three Embarcadero Center
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING STEEM: DOS
SOFTWARE: FASTESD for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/784,747
FILING DATE: L6-JAN-1997
CLASSIFICATION DATA:
APPLICATION NUMBER:
FILING DATE: EXAMPLED:
FILING DATE: FILING DATA:
ATORNEY/AGENT INFORMATION:
NAME: KANDAMOLO, ANGREW A
RECISTRATION NUMBER: 40,690
REFERENCE/DOCKET NUMBER: 20411-708
TELECOMMULCATION INFORMATION:
TELECOMMULCATION INFORMATION:
TELECOMMULCATION INFORMATION:
TELECOMMULCATION INFORMATION:
TELEFAX: 415-393-2286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1146 TTTTTTTTTGG 1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 TTTTINTTITEG 15
                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 92.3
Matches 12; Conservative
                              COMPUTER: IBM Com
                                                                                                                                                                                                                                                                                                                                                                                                              linear
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US-08-784-747-2
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OTHER INFORMATION: Exemplary oligonucleotide primer used in sequence assembly proc.; NAME/KEY: misc_feature; LOCATION: (8)...(8)

COCATION: (8)...(8)

US-09-403-778-9
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Sequence 9, Application US/09409778

Sequence 9, Application US/09409778

Sequence 9, Application US/09409778

GENERAL INFORMATION:
APPLICANT: Ford, John
APPLICANT: Pord, John
TITLE OF INVENTION: A NOVEL CHEMOKINE RECEPTOR OBTAINED FROM TITLE OF INVENTION: A NOVEL CHEMOKINE RECEPTOR OBTAINED FROM TITLE OF INVENTION: A NOVEL CHEMOKINE RECEPTOR OBTAINED FROM TITLE OF INVENTION: A NOVEL CHEMOKINE RECEPTOR OBTAINED FROM TITLE OF INVENTION: A NOVEL CHEMOKINE RECEPTOR OBTAINED FROM TITLE OF INVENTION NUMBER: US/09/409,778

CURRENT APPLICATION NUMBER: 1999-09-22

PRIOR PILING DATE: 1999-06-29

PRIOR APPLICATION NUMBER: US/09/236,166

PRIOR APPLICATION NUMBER: US/09/236,166

PRIOR PILING DATE: 1999-01-22

PRIOR PILING DATE: 1999-06-26

SEQ ID NOS: 25

SEQ ID NOS: 25

SEQ ID NOS: 25

LENGTHE: FASTSEQ FOR WINDOWS VERSION 3.0

SEQ ID NO 9
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| Sequence 10, Application US/09409778 |
| Sequence 10, Application US/09409778 |
| Patent No. 6472173 |
| General Information: |
| APPLICANT: Ford, John |
| APPLICANT: Ford, George |
| TITLE OF INVENTION: A CHARACTION CONTAINED FROM |
| TITLE OF INVENTION: A CHARACTION (Inow 28110/36057B) |
| CURRENT APPLICATION UNDERS: US/09/409,778 |
| CURRENT FILING DATE: 1999-06-22 |
| PRIOR PILING DATE: PCT/US99/12829 |
| PRIOR FILING DATE: US 09/236,166 |
| PRIOR APPLICATION NUMBER: US 09/236,166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                        Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.9%; Score 12; DB 1; Length 15; 92.3%; Pred. No. 1.8e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                           1; Indels
                                                                                                                                                                                                                   Score 12; DB 1; Dred. No. 1.8e+02; 0; Mismatches 1,
                                                                                                                                                                                                                   0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                           1146 TTTTTCTTTTGG 1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1146 TTTTTCTTTTGG 1158
                                                                                                                                                                                                       Query Match
Best Local Similarity 92.39
Matches 12; Conservative
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 92.33
Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                13 TTTTTNTTTTGG 1
                                                                                                             single
                                                                                 TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                       linear
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Gaps

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ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/08/882,649A
FILING DATE: 25-Jun-1997
CLASSIFICATION: 435-006.000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tran, Huu M.
Matsuzaki, Hajime
McGall, Glenn H.
TITLE OF INVENTION: NUCLEIC ACID ANALYSIS TECHNIQUES
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Joe Liebeschuetz
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Liebeschuetz, Joe
REGISTRATION NUMBER: 37.505
REFERENCE/DOCKET NUMBER: 018547-019410US
TELECOMMUNICATION INFORMATION:
                              Pred. No. 2e+02;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
0.9%; Score 12; DB 1;
Best Local Similarity 81.2%; Pred. No. 2e+02;
Matches 13; Conservative 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/010,471
FILING DATE: 23-JAN-1996
APPLICATION NUMBER: US 60/035,170
FILING DATE: 09-JAN-1997
APPLICATION NUMBER: PCT/US97/01603
FILING DATE: 22-JAN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ix) Features:
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (415) 576-0200
INFORMATION FOR SEC ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                       Sequence 8, Application US/08882649A
Patent No. 6344316
GENERAL INFORMATION:
APPLICANT: Lockhart, David J.
Chee, Mark
Gunderson, Kevin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chaogiang, Lai
Wodicka, Lisa
Cronin, Maureen T.
100.001;
14 :80.001
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MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1141 GCCTTTTTTTTTT 1156
                                                                                                                      576 GCAGGCCCTCCG 587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lee, Danny
                    Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                   15 GCAGGCCCTCCG 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
                                                                                                                                                                                                                                                                              RESULT 446 '
US-08-882-649A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-882-649A-8
                                                                                                                                                                                   a
                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.9%; Score 12; DB 1; Length 15; Best Local Similarity 92.3%; Pred. No. 1.88+02; Matches 12; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Birch, Stewart, Kolasch & Birch, Lide STREET: 8110 Gatehouse Road, Suite 500 East CITY: Fallo Church STATE: Virginia COUNTRY: U.S.A.

ZIB: 22042 COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk COMPUTER: Floppy disk COMPUTER: Ploppy disk COMPUTER: Ploppy disk COMPUTER: Ploppy disk COMPUTER: Plan PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION NUMBER: US/08/232,087A FILING DATE: TRING DATE: TRING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Stein, Harald
APPLICANT: Drkop, Horst
APPLICANT: Latza, Ute
TITLE OF INVENTION: Lymphoid CD30-Antigen
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSE: Birch, Stewart, Kolasch & Bir
                                                                                                               SEQ ID NOS: 25
FastSEQ for Windows Version 3.0
          PRIOR FILING DATE: 1999-01-22
PRIOR APPLICATION NUMBER: US 09/106,800
PRIOR FILING DATE: 1998-06-26
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FASTSEQ for Windows Version 3.
SEQ ID NO 10
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-232-087A-5/c; Sequence 5, Application US/08232087A; Patent No. 5866372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 08-SEP-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MUTCHY Jr., Gerald M.
REGISTRATION NUMBER: 28,977
REPROCE/DOCKET NUMBER: 756-
TELEPHONE: (703) 205-8000
TELEPAX: (703) 205-8050
                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1146 TTTTTTTTTGG 1158
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INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13 TTTTTTTGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
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ORIGINAL SOURCE:
ORGANISM: HOM
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Gaps

0.9%; Score 12; DB 1; Length 16;

Query Match

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OTHER INFORMATION: Description of Artificial Sequence:Designed OTHER INFORMATION: oligonucleotide based on the amino acid sequence OTHER INFORMATION: represented in SEQ ID NO.4.
           EARLIER APPLICATION NUMBER: 10-234769 JAPAN
                                    EARLIER FILING DATE: 1998-08-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                    629 AGCICCAGGAGCICIG 644
                                                                                                                                         TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 AGCTYCASRASCCVTG 17
                                                NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 14
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                        10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: CDNA HYDOTHRTICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   unknown
                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FRAGMENT TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-984-709A-45
                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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66.7%; Pred. No. 2.3e+02;
ive 4; Mismatches 0; Indels
                                                                                                  Sequence 649, Application US/08758306
Fatent No. 5807743
GENERAL INFORMATION:
APPLICANT: Stinchcomb, Dan T.
APPLICANT: MCSWiggen, James A.
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES
TITLE OF INVENTION: TREATMENT OF DISEASES
TITLE OF INVENTION: INTERLEUKIN-2 RECEPTOR
TITLE OF INVENTION: MASCOLATED WITH
TITLE OF INVENTION: GAMMA-CHAIN EXPRESSION
NUMBER OF SEQUENCES: 1379
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Los Angeles
CONTANT: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 14, Application US/09328501A
Patent No. 6258591
GENERAL INFORMATION:
APPLICANT: ITO, Makoto
ITILE OF INVENTION: Ceramidase Gene
FILE REFERENCE: 1422-0377p
CURRENT APPLICATION NUMBER: US/09/328,501A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: CALLIOTHIA
COUNTRY: CALLIOTHIA
COUNTRY: CALLIOTHIA
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM COMPALIA
COMPUTER: IBM COMPALIA
COMPUTER: IBM COMPALIA
COMPUTER: IBM COMPALIA
COMPUTER: IBM P.C. DOS 5.0
SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/758,306
FILING DATE: December 3, 1996
CLASSIETCATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
FILING APPLICATION NUMBER:
FILING APPLICATION NUMBER:
RECISTRATION NUMBER:
ATTORNEY AGENT INFORMATION:
NAME: WATBURG, RICHARD J.
REFERRATOR WUMBER: 32,327
REFERRATOR MUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBES: 23
TELECOMMUNICATION INFORMATION: (213) 489-160
TELEFRAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212)
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 64
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
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Best Local Similarity 66.,
Associated 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       83 AATAGCAGTTCT 94
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TOPOLOGY:
US-08-758-306-649
                                                                                  US-08-758-306-649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-328-501-14
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Gaps
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APPLICANT: Standerman, Kenneth A.
APPLICANT: Standerman, Kenneth A.
APPLICANT: Harbold, Michel M.
APPLICANT: Harbold, Michel M.
TITLE OF INVENTION: HUMAN CALCIUM CHANNEL COMPOSITIONS AND
TITLE OF INVENTION: METHODS
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White & McAuliffe
STREET: 4250 Executive Square, Suite 700
CITY: La Jolla
STATE: California
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:

NAME: Seidman, Stephanie L.

REGISTRATION NUMBER: 33,779

REFERENCE/DOCKET UNDHER: 24735-9815 (formerly 6362-9815)

TELECOMMUNICATION INFORMATION:

TELEPHONE: (619) 450-8400

TELEPAX: (619) 587-5360

INFORMATION FOR SEQ ID NO: 45:

SEQUENCE CHARACTERISTICS:

LENGTH: 17 base pairs

TYPE: nucleic acid

STRANDEDNESS: single
                                                                                               0.9%; Score 12; DB 1; Length 17; 62.5%; Pred. No. 2.38+02; ive 5; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
FEATURE:

OTHER INFORMATION: any n or Xaa = Unknown US-09-328-501-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.9%; Score 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/984,709A
FILING DATE: 02-DEC-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                     RESULT 449
US-08-984-709A-45
Sequence 45, Application US/08984709A
Patent No. 6320032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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RESULT 450

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Gaps
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00.0%; Pred. No. 2.3e+02;
ve 0; Mismatches 0; Indels
                                              APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Estinchcomb, Dan T.
APPLICANT: Estinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF SEQUENCES: 6502
CORRESPONDENCE ADDRESS:
ADDRESSER: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Laboratory of Molecular Biophotonics
ITLE OF INVENTION: Method for gene analysis
FILE PERERENCE: 400595
CURRENT APPLICATION NUMBER: US/09/537,720B
CURRENT FILING DATE: 2000-03-30
PRIOR APPLICATION NUMBER: JP12/22630
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin version 3.1
SEQ ID NO 15
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0%; Pred. No. z...
Matches 12; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
SOFTWARE: WORD PERCET 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: OCCODER: 26, 1995
ATTORNEY, AGENT INFORMATION:
NAME: WALDING, RICHARD J.
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                       CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 218/064
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 21.
TELECOMMUNICATION: 1NFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                              Pavco, Pamela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              909 CCTGGTCCTAAA 920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 17 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TOPOLOGY: linear
US-08-584-040-7538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-537-720B-15/c
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                            APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 452
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                              Gaps
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                         Indels
                                                                                                                                                                                                                                                                                 APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Stinchomb, Dan T.
APPLICANT: Escobedo, Jaine
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: GROWTH FACTOR
75.0%; Pred. No. 2.3e+02; tive 2; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTITE T. U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYZETHE: IBM P.C. DOS 5.0
SOFTWARE: WORD Perfect 5.1
CURRENT APPLICATION DATA:
FILING DATE: January 11, 1996
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: WARDLEY RICHARD J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                  Sequence 1844, Application US/08584040
Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-584-040-7538/c; Sequence 7538, Application US/08584040; Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 1844:
SEQUENCE CHARACTERISTICS:
                                                               323 ACCTGCATCATCTGG 338
                                                                                           1 |||:||:| |||||
1 AACTGYATYACCCTGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
Best Local Similarity 75.0
Matches 12; Conservative
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Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 ccrédrecrasa 6
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STATE: California
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                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
                                                                                                                                                                                                  US-08-584-040-1844/C
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RESULT 451

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629 AGCICCAGGAGCICIG 644
                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                          62.5%;
                                                                                                                                                                                                                                                                                                                                            10; Conservative
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US-09-371-772B-389
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Best Local Similarity
                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 10; Conserv
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US-09-371-772B-4638/c
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Patent No. 6433155
GENERAL INFORMATION
APPLICANT: Umansky, Samuil
APPLICANT: Welkonyan, Hovsep
TITLE OF INVENTION: APOPTOSIS-RELATED PEPTIDES ENCODED THEREBY AND
TITLE OF INVENTION: METHODS OF USE THEREOF
TITLE OF INVENTION: METHODS OF USE THEREOF
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSE: MORISSON & FOERSTER
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: CA
COUNTRY: USA
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80.0%; Pred. No. 2.38+02;
tive 1; Mismatches 2; Indels
                                                                                                                                      0.9%; Score 12; DB 1; Length 17;
100.0%; Pred. No. 2.3e+02;
Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA

ZIP: 94304-1018

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/937,067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILLENGHEN DALE:
CLASSIFICATION:
NAME: Lebnhardt, Susan K.
NAME: Lebnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 23647-20018.00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEPHONE: (650) 494-0792
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: mucleic acid
STRANDENBES: single
                FEATURE:
COTHER INFORMATION: Predicted Sequence
NAME/KEY: misc feature
CTHER INFORMATION: Part of the template
US-09-537-720B-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 14, Application US/09777710A; Patent No. 6489117; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1144 TITITITITIGG 1158
ORGANISM: Artificial Sequence
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TTTTTTTTTTTNS 17
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Best Local Similarity 100.
Matches 12; Conservative
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Best Local Similarity 80.0
Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 454
US-09-777-710A-14
                                                                                                                                                                                                                                                                                                                                  US-08-937-067-17
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APPLICANT: Pavco, Pam
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcowb, Dan
APPLICANT: Stinchcowb, Dan
APPLICANT: Stinchcowb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
FILE OF INVENTION: McWels of Vascular Endothelial Growth Factor Receptor
FILE OF INVENTION: McWels: US/09/371,772B
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PLIANG DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR PLIANG DATE: 1995-61-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEQ ID NO 389
LENGTH: 17
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APPLICANT: McSwiggen, Jim
APPLICANT: Stinchomb, Dan
APPLICANT: Escobedo, Jaime
IITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Designed oligonucleotide based on the amino acid sequence is OTHER INFORMATION: represented in SEQ ID NO:4 US-09-777-7108-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 12; DB 1; I
52.5%; Pred. No. 2.3e+02;
ive 5; Mismatches 1;
APPLICANT: OKINO, NO. 64891170mu et al.
TITLE OF INVENTION: CERAMIDASE GENE
FILE REFERENCE: 1422-0458P
CURRENT APPLICATION NUMBER: US/09/777,710A
CURRENT FILING DATE: 2001-02-07
NUMBER OF SEQ ID NOS: 18
SEQ ID NO 14
LENGTWARE: Patentin version 3.0
SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Pred. .v.
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18 Sequence 389, Application US/09371772B

Patent No. 6566127

GENERAL INFORMATION:

APPLICANT: Ribozyme_Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 4638, Application US/09371772B
Patent No. 6566120
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
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Gaps
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                                                                                                                                                                                     0.9%; Score 12; DB 1; Length 17; 92.3%; Pred. No. 2.38+02; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 1930405
CLASSIFICATION: 536
ATTORNEY AGENT INFORMATION:
REFERENCE/DOCKET NUMBER: 20276/SBI36848/US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: MCLEAN, MICHAEL J.
APPLICANT: HOLLAND, DAVID
APPLICANT: GARWAN, ANDREW J.
APPLICANT: SHEPPARD, ROBERT C.
TITLE OF INVENTION: SYNTHESIS OF OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.9%; Score 11.8; DB 1;
Best Local Similarity 86.7%; Pred. No. 1.9e+02;
Matches 13; Conservative 0; Mismatches 2;
       /note= "5-methylcytosine"
                                                                                               /mod_base= OTHER
/note= "1,3-propanediol"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: CUSHMAN, DARBY & CUSHMAN STREET: 1100 NEW YORK AVENUE, N.W. CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 459
US-08-127-954-50
; Sequence 50, Application US/08127954
; Parent No. 5451512
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2, Application US/08041599
Patent No. 5393877
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1145 TTTTTTTTTTGGA 1159
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INPORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                               1144 TTTTTTTTTT 1156
                            FEATURE:
NAME/KEY: modified_base
                                                                                                                                                                                     Query Match 0.95
Best Local Similarity 92.35
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                          1 TTTTTTTTTTT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleic acid
EDNESS: single
                                                                       LOCATION: 17
CTHER INFORMATION:
CTHER INFORMATION:
PCT-US91-03680-7
       OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: WASHIN STATE: D.C. COUNTRY: U.S ZIP: 20005
                                                                                                                                                                                                                                                                                                                                                                                           RESULT 458
US-08-041-599-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-041-599-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 7, Application PC/TUS9101680
GENERAL INFORMATION:
APPLICANT: Matteucci, Mark D.
APPLICANT: Krawczyk, Steven
TITLE OF INVENTION: SEQUENCE-SPECIFIC NONPHOTOACTIVATED
TITLE OF INVENTION: CROSSIINKING AGENTS WHICH BIND TO THE MAJOR GROOVE OF
NUMBER OF SEQUENCES: 158
                                                                                                                                                                                                                                                                                                                                                                                             ó
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                                                                                                                                                                                                                                                                                                                                             DB 1; Length 17; 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: ...
ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US91/03680
FILING DATE: 19910524
CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: MULESHIGE Kate H.
REGISTRATION NUMBER: 29,959
REPRENCE/DOCKET NUMBER: 4610-0011.40
TELECHONGUINGATION INFORMATION:
TELEPHONE: 415-327-2951
TELEPHONE: 415-327-2951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "N4,N4-ethanocytosine"
                CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT PILLING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILLING DATE: 1995-10-26
PRIOR FILLING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                      Query Match 0.9%; Score 12; DB Best Local Similarity 100.0%; Pred. No. 2.3 Matches 12; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSE: Morrison & Foerster
STREET: 845 Middlefield Road, Suite 200
CITY: Menlo Park
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: /mod_base= OTHER OTHER INFORMATION: /note= "N4, N4-etherthe:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: 14 OTHER INFORMATION: /mod_base= OTHER
FILE REFERENCE: MBHB00,876-J (237/198)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELERAX: 415-327-2951
TELEX: 706141
INFORMATION FOR SEQ ID NO: 7
SEQUENCE CHARACTERISTICS:
LENOTH: 17 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: modified_base LOCATION: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12 ceregrecraaa 1
                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
                                                                                                                                                                                                                                                                                              US-09-371-772B-4638
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; Sequence 8, Application US/08276099A
; Patent No. 5591825
; GENERAL INFORMATION:
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               07-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 910 277299
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 15 base pairs TYPE: nucleic acid STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: linear; MOLECULE TYPE: cDNA US-08-337-025-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
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US-08-276-099A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-276-099A-8
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                       APPLICANT: Apple, Raymond J.
APPLICANT: Bugawan, Tedorica L.
APPLICANT: Bugawan, Tedorica L.
APPLICANT: Brlich, Henry A.
TITLE OF INVENTION: Locus DNA Typing
NUMBER OF SECUENCES: 173
CORRESPONDENCE ADDRESS:
ADDRESSE: Hoffmann-La Roche Inc.
STREET: 340 Kingsland Street
CITY: Nutley
STREET: New Jersey
COUNTRY: 0.3.A.
ZIP: 071.0-1199
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: BLE PC COMPUTER: PLOSS MS-DOS
CORRESPENT OF STREET: PO-DOS/MS-DOS
CORRESPENTED: PLOSTEM: PC-DOS/MS-DOS
CORRESPENTED: PC-DOS/MS-D
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86.7%; Pred. No. 1.9e+02;
tive 0; Mismatches 2; Indels
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Faquence 1, Application US/08337025

GENERAL INFORMATION:
APPLICANT: MCLEAN, MICHAEL J.
APPLICANT: HOLLAND, DAVID
APPLICANT: GARMAN, ANDREW J.
TITLE OF INTERPRAD, ROBERT C.
TITLE OF SEQUENCES: 2
CORRESPONDENCES: 2
CORRESPONDENCES: CUSHWAN, DARBY & CUSHWAN
STREET: 1100 NEW YORK AVENUE, N.W.
CITY: WASHINGTON
STATE: D.C.
COUNTRY: U.S.A.
CANTENT: 20005
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MEDIUM TYPE: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/WS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/337,025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/127,954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICALLA. ...
FILLING DATE:
CLASSIFICATION: 436
ATTORNEY/AGENT INFORMATION:
NAME: PELTY, DOUGLAS.
REGISTRATION NUMBER: 35,321
REFERENCE/DOCKET NUMBER: 35,321
TELECOMUNICATION INFORMATION:
TELEPHONE: (510) 814-2974
TELEPHONE: (510) 814-2977
; INPORMATION FOR SEQ ID NO: 50; SEQUENCE CHARACTERISTICS;
LENGTH: 15 base pairs
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-127-954-50
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Best Local Similarity 86.7°
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleic acid
EDNESS: single
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-337-025-2
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APPLICANT: MCALIGHT, Steven L
APPLICANT: Hou, Jinzhao
TITLE OF INVENTION: INTERLETKIN-4 SIGNAL TRANSDUCERS AND
TITLE OF INVENTION: BINDING ASSAYS
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLERR, HOHBACH, TEST, ALBRITTON & HERBERT
STRET: 4 Embarcadero Center, Suite 3400
CITY: 5an Francisco
STATE: California
COUWTRY: USA
ZIP: 194111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: RIDAPY disk
COMPUTER: DELAFIL Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 18/08/276,099A
FILING DATE: 15-JUL-1994
CLASSIFICATION NUMBER: 36,627
REGISTRANTION NUMBER: 36,627
RELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 398-3249
TELEPRAN: 610,077-304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.9%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 1.9e+02; Matches 13; Conservative 0; Mismatches 2; Indels
CLASSIFICATION: 536
PIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/041,599
FILING DATE: 05-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: BIRD, DONALD J.
REGISTRATION NUMBER: 25,323
REFERENCE/DOCKET NUMBER: 202706/SBI36848/US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-803-44
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
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Gaps
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                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE PORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM COMPACTION
COMPUTER: IBM COMPACTION
COMPUTER: IBM COMPACTION
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/291,932A
CLASSIFICATION S14
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
APPLICATION DATA: described below:
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION
AND ATTORNEY PROFESSION DESCRIPTION
ATTORNEY PROFESSION
ATTORNEY PROFESSION DESCRIPTION
ATTORNEY PROFESSION
ATTORNEY PR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.9%; Score 11.8; DB 1;
Best Local Similarity 86.7%; Pred. No. 1.9e+02;
Matches 13; Conservative 0; Mismatches 2;
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APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: NP-KB
NUMBER OF SEQUENCES: 830
CORRESPONDENCE ADDRESS:
      DISEASES OR CONDITIONS
                                 RELATED TO LEVELS OF NF-KB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 464
US-08-291-932A-378/c
'Sequence 378, Application US/08291932A
'Patent No. 5558780
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                       ADDRESSEE: Lyon E Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 200
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMPOST. (213) 489-10-1 TELEPAOR (213) 955-0440
TELEX. 67-3510
INFORMATION FOR SEQ ID NO: 33
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6 GGCAGTTGAGGTGGA 20
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TITLE OF INVENTION: DISE
TITLE OF INVENTION: RELA
TITLE OF INVENTION: NF-K
NUMBER OF SEQUENCES: 830
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY:
US-08-291-932A-33
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                                                                      Gaps
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                                                                      0
Query Match

0.9%; Score 11.8; DB 1; Length 15;

Best Local Similarity 86.7%; Pred. No. 1.9e+02;

Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.9%; Score 11.8; DB 1; Length 15; 53.3%; Pred. No. 1.9e+02; tive 5; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 463
US-08-291-932A-33/C
; Sequence 33, Application US/08291932A
; Patent No. 5658780
; GENERAL INFORMATION:
; APPLICANT: Stinchcomb, Dan T.
; APPLICANT: Dxaper, Kenneth G.
; APPLICANT: McSwiggen, James
; TITLE OF INVENTION: RIBOZYME TREATMENT OF
                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT:
APPLICANT:
TITLE OF INVENTION:
METHOD AND REAGENT FOR
TITLE OF INVENTION: INHIBITING HEPATITIS C
TITLE OF INVENTION: VIRUS REPLICATION
NUMBER OF SEQUENCES: 497
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSES: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STRATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: BACARGE
COMPUTER: BM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/182,968A
FILING DATE: 13-OANUARY-1994
FRIOR APPLICATION NUMBER: 07/882,888
FRIOR APPLICATION NUMBER: 07/882,888
FRIOR APPLICATION NUMBER: 205/277
REFERENCE/DOCKET NUMBER: 205/277
REPERENCE/DOCKET NUMBER: 205/277
TELECOMMUNICATION INFORMATION:
TELESK: 67-3510
TELEBEX: 67-3510
TELEBEX: 67-3510
TINPORMATION FOR SEG ID NO: 201:
                                                                                                                                                                                                                                                                                                                                                      ; Sequence 201, Application US/08182968A; Patent No. 5610054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1190 TGAGTGTTGGACCTT 1204
                                                                                                                               835 AAGCTTTCAGATGGG 849
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Best Local Similarity 53.3%
Best Local 8; Conservative
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STRANDEDNESS: sing];
TOPOLOGY: linear
US-08-182-968A-201
                                                                                                                                                                                                                                                                                                                           US-08-182-968A-201
                                                                                                                                                                                                                                                                                   RESULT 462
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.9%; Score 11.8; DB 1; Length 1 Best Local Similarity 20.0%; Pred. No. 1.9e+02; Matches 3; Conservative 10; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 606, Application US/08334847
Patent No. 5693532
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: McSwiggen, James
APPLICANT: PavCo, Pam
APPLICANT: PavCo, Pam
APPLICANT: PovOol, Tod
ITTLE OF INVENTION: METHOD AND REAGENT FOR
ITTLE OF INVENTION: SYNCYTIAL VIRUS
NUMBER OF INVENTION: SYNCYTIAL VIRUS
NUMBER OF INVENTION: SYNCYTIAL VIRUS
NUMBER OF INVENTION: LYON & LYON
NUMBER OF SEQUENCES: 909
CORRESPONDENCE ADDRESS:
ADDRESSEE: LYON & LYON
FILE OF INVENTION: SYNCYTIAL VIRUS
NUMBER OF SEQUENCES: 909
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 30071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/334,847
                  FILING DATE: No. 5693532ember 4, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: No. 5693532ember 4, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 209/032
TELECOMUNICATION INFORMATION:
TELEPHONE: (213) 495-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                           209/032
                                                         ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        909
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                         LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15 base pairs
                                      PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY:
US-08-334-847-570
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-334-847-606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.9%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 1.9e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Indels
CCOUNTRY: U.S.A.

ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/291,932A
FILING DATE: August 15, 1994
CLASSIFICATION DATA: Gescribed below:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
PRIOR APPLICATION DATA: described below:
PRIOR APPLICATION NUMBER: 09/287,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 20/987,132
FILING DATE: December 7, 1992
ATTORNEY AGBNT INFORMATION:
REGISTRATION NUMBER: 20/987,137
FELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-160
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
TELEX: 67-3510
INPORMATION FOR SEQ ID NO: 378:
LENGTH: 15 base pairs
TTRANSENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Pavco, Pam
APPLICANT: Woolf, Tod
ITLE OF INVENTION: METHOD AND REAGENT FOR
ITLE OF INVENTION: INHIBITING RESPIRATORY
ITLE OF INVENTION: SYNCTIAL VIRUS
NUMBER OF SEQUENCES: 909
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: SULLE 4/00
CITY: LOS Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-206
COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/334,847
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 570, Application US/08334847 Patent No. 5693532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          528 GGAGGAGCAGCTGGG 542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15 GGAGCAGGAGCTGGG 1
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
TOPOLOGY:
US-08-291-932A-378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 465
US-08-334-847-570
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Sequence 142, Application US/08363240A Patent No. 5705388
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                                                                                                                    Query Match 0.9%; Score 11.8; DB 1; Length 15; Best Local Similarity 46.7%; Pred. No. 1.9e+02; Matches 7; Conservative 6; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                      Sequence 631, Application US/08334847
Fatent No. 5693532
GENERAL INPORMATION:
APPLICANT: MCSWiggen, James
APPLICANT: Draper, Kenneth
APPLICANT: PavCo, Pam
APPLICANT: PavCo, Pam
APPLICANT: PavCo, Pam
APPLICANT: Moolf, Tod
TITLE OF INVENTION: INHIBITING RESPIRATORY:
TITLE OF INVENTION: SYNCYTIAL VIRUS
NUMBER OF SEQUENCES: 909
CORRESPONDENCE: 2009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Los Aqueles
STATE: California
COUNTRY: U.S.A.

ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 18M Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/334,847
FILING DATE: No. 5693532ember 4, 1994
PRIOR APPLICATION NUMBER:
FILING DATE:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: WALDLUG, RIChard U.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/032
TELECHMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                               442 AAGTTGCTGAAGTTT 456
                                                                                                                                                                                                                                         1 AAGUUGUUGAGGUUU 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 60.09
Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-334-847-606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-334-847-631
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Gaps
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             APPLICANT: COLULE, LAITY
APPLICANT: McSwiggen, James
APPLICANT: Bisgater, Charles
APPLICANT: Bisgater, Charles
APPLICANT: Pape, Michael
TITLE OF INVENTION: PREVENTION INHIBITION OF
TITLE OF INVENTION: PREVENTION, INHIBITION OF
TITLE OF INVENTION: PREVENSION AND REGRESSION
TITLE OF INVENTION: OF VASCULAR DISEASES
NUMBER OF SEQUENCES: 1243
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: 633 West Fifth Street
CITY: Los Angeles
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Couture, Larry
APPLICANT: MCSWiggen, James
APPLICANT: Bisgaier, Charles
APPLICANT: Pape, Michael
ITILE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: PREVENTION, INHIBITION OF
TITLE OF INVENTION: OF VASCULAR DISEASESION
TITLE OF INVENTION: OF VASCULAR DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage COMPUTER: 1EM COMPATIBLE OPERATING SYSTEM: IEM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/363,240A FILING DATE: December 23, 1994 PRIOR APPLICATION NUMBER: US/08/363,240A PRIOR APPLICATION NUMBER: US/08/363,240A PRIOR APPLICATION NUMBER: US/08/363,240A PRIOR APPLICATION NUMBER: 33, 137 REFERENCE/DOCKET NUMBER: 32, 327 REFERENCE/DOCKET NUMBER: 210/096 TELECOMMUNICATION INFORMATION: TELEFRONE; (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-363-240A-541/c
; Sequence 541, Application US/08363240A
; Patent No. 5705388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               307 GGGCTGCAACTCCA 321
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 80.05
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A. ZIP: 90071
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
TOPOLOGY: ]
US-08-363-240A-142
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; Sequence 8. Application US/08781890
; Sequence 8. Application US/08781890
; Patent No. 5710266
; Patent No. 5710266
; Patent No. 5710266
; APPLICANT: McKnight, Steven L
; APPLICANT: McKnight, Steven L
; APPLICANT: McKnight, Steven L
; TITLE OF INVENTION: INTERLEUKIN-4 SIGNAL TRANSDUCERS AND
TITLE OF INVENTION: INTERLEUKIN-4 SIGNAL TRANSDUCERS AND
TITLE OF INVENTION: BINDING ASSAYS
; NUMBER OF SEQUENCES:
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 11.8; DB 1; Length 15; Pred. No. 1.9e+02; 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CUDNIES: CALIFORNIA

ZIP: 94111-4187

COUNTER: USA

COMPUTER READABLE FORM:
MEDIUW TYEE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHIN PC-DOS/MS-DOS
SOFTWARE: PATCHIN DATA:
APPLICATION NUMBER: US/08/781,890
FILING APPLICATION 536
PRIOR APPLICATION NUMBER: US/08/781,890
FILING APPLICATION 1536
PRIOR APPLICATION NUMBER: US/08/76,099
FILING APPLICATION NUMBER: US/08/76,099
FILING APPLICATION NUMBER: A-59461-1/RAO
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: 3-59451-1/RAO
TELECOMMUNICATION INFORMATION:
TELEFRAK: (415) 781-1989
TELEFRAK: (415) 781-1989
                    US/08/363,240A
                                              FILING DATE: December 23, 1994 PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                     FILING DATE:
ATORNEY/AGENT INFORMATION:
NAME: WAZDUZG, RICHARD
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 210/C
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFA: (7-3510
INFORMATION FOR SEQ ID NO: 658:
SEQUIENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: mucle; acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GGGCAGGCAGTTGAG 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15 GCCCAGGGAGTTGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 86.7%
Watches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 910 277,939-324
TELEX: 910 277,939-324
INFORMATION FOR SEQ 120 NO: SEQUENCE CHARACTERISTICS: LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: STRANDEDNESS: SIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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STRANDEDNESS:
TOPOLOGY: line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-363-240A-658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 471
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86.7%; Pred. No. 1.9e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GEOUGH 658/C

Sequence 658, Application US/08363240A

Patent No. 5705388

Sequence 658, Application US/08363240A

Patent No. 5705388

MENERAL INFORMATION:
APPLICANT: McSwigaen, James
APPLICANT: Bisgaier, Charles
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: PREVENTION, INHIBITION OF
TITLE OF INVENTION: PROCRESSION AND REGRESSION
TITLE OF INVENTION: PROCRESSION
TITLE OF INVENTION: PROCRESSION
TITLE OF INVENTION: PROCRESSION
TITLE OF INVENTION: PROCRESSION
TITLE OF INVENTION:
TITLE OF INVENTION: PROCRESSION
TITLE OF INVENTION:
TIT
                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 8torage
COMPUTER: IRM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PETFECT 5.1
CURRENT PAPPLICATION DATA:
APPLICATION NUMBER: US/08/363,240A
FILING DATE: December 23, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/363,240A
FILING DATE: APPLICATION DATA:
APPLICATION NUMBER: 23, 1994
REFERENCE/AGENT INPORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 210/096
TELECOMMUNICATION INPORMATION:
TELECHNONE: (213) 489-1600
TELEFRAX: (213) 955-0440
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       924 GATGGCAGATCTGGA 938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 5-
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15 GGTGGCTGATCTGGA 1
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Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-363-240A-541
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Db

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Best Local Similarity 86.7%; Pred. No. 1.9e+02;
Matches 13; Conservative 0; Mismatches 2;
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US-08-292-620A-74
                                                                                                                                                                                           RESULT 473
US-08-292-620A-74
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                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Warren, Gregory W
APPLICANT: Koziel, Michael G
APPLICANT: Mullins, Martha A
APPLICANT: Mullins, Martha A
APPLICANT: Desai, Gordon J
APPLICANT: Carr, Brian
APPLICANT: Carr, Brian
APPLICANT: Costichka, Nalini M
APPLICANT: Estruch, Nalini M
APPLICANT: Estruch, Juan J
APPLICANT: Estruch, Juan J
APPLICANT: No. 5770696el Pesticidal Proteins and Strains
NUMBER OF SEQUENCES: 50
                                                                                                                        ;
0
                                                                      0.9%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:

CLASSIFICATION DATA:

PRIOR APPLICATION NUMBER: US 08/314,594

PRIOR APPLICATION NUMBER: US 08/314,594

PRIOR APPLICATION NUMBER: US 08/218,018

PRIOR APPLICATION NUMBER: US 08/218,018

PRIOR APPLICATION NUMBER: US 08/218,018

PRIOR APPLICATION NUMBER: US 08/037,057

PRIOR APPLICATION NUMBER: US 08/037,057

PRIOR APPLICATION NUMBER: P40,403

REFINENCE/DOCKET NUMBER: P1-41-6892

TELECOMMUNICATION INFORMATION:

SEQUENCE CHARACTERISTICS:

LENGTH: 15 base pairs

TYPE: nucleic acid

TYPE: nucleic acid
                                                                                                                     2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               other nucleic acid
/desc = "reverse primer used to make
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 10532

ZIP: 10532

COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30B
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/471,033
                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSER: CIBA-GETGY Corporation
STREET: 7 Skyline Drive
CITY: Hawthorne
                                                                                                                                                                                                                                                                                     RESULT 472
US-08-471-033-34/c
Sequence 34, Application US/08471033
; Patent No. 5770696
                                                                                                                                                                     835 AAGCTTTCAGATGGG 849
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                                                                                                                   Matches 13; Conservative
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                                                                                               Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Warren
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: NY
COUNTRY: USZ
ZIP: 10532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-471-033-34
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US-08-781-890-8
                                                                        Query Match
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0.9%; Score 11.8; DB 1; Length 15;

Query Match

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Gaps
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Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Los Angeles
CITY: Los Angeles
COUNTRX: Los Angeles
COUNTRX: U.S.A.
ZIP: 30071-2066
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 1BM COMPATIBLE
COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
COMPUTER: Nord Perfect 5.1
CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/292,620A
FILING DATE: August 17, 1994
CLASSIPICATION DATA: including application
PRIOR APPLICATION DATA: described below:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: 1994
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEPHONE (23) 489-1660
TELEPHONE (23) 485-1660
                                                                                                                                                                                                       Sequence 74, Application US/08292620A

Patent No. 5837542

GENERAL INFORMATION:
INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: Games McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
APPLICANT: HENDER TREATMENT OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: NULECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Sulfe 4700
                                                826 ATGCAGCTGAAGCTT 840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (412),
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 74
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            908 CCCTGGTCCTAAAGG 922
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 11; Conserv
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ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 51-0196
COMPUTER: 1BM COMPATIBLE
COMPUTER: 1BM COMPATIBLE
COMPUTER: 1BM COMPATIBLE
COMPUTER: 1BM COMPATIBLE
COMPUTER: 1BM P.C. DOS 5.0
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 10, 1994
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: 08/008/895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 08/008/895
FILING DATE: DECEMBER 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGIERRANCE/DOCKET NUMBER: 20/149
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Suean Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: RELATED TO LEVELS OF
                                                Kenneth G. Draper
NVENTION: RIBOZYME TREATMENT OF
NVENTION: DISEASES OR CONDITIONS
NVENTION: RELATED TO LEVELS OF
NVENTION: INTRACELLULAR ADHESION
NVENTION: MOLECULE-1 (I-CAM-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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US-08-292-620A-656/c
; Sequence 656, Application US/08292620A
; Patent No. 5837542
                                                                   TITLE OF INVENTION: REBOZYME TREAT
TITLE OF INVENTION: RELATED TO LEY
TITLE OF INVENTION: RELATED TO LEY
TITLE OF INVENTION: RELATED TO LEY
TITLE OF INVENTION: MOLECULE-1 (I-
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: LYON & LYON
STREET: 633 West Fifth Street
STREET: SULF 4700
CITY: LOS Angeles
STREET: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1260 CCAGGTTGAGGCCCT 1274
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
James McSwiggen
Sean Sullivan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.9
Best Local Similarity 86.7
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-292-620A-393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX:
                                                  APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLIANG DATE: AUGUET.

CLASSIFICATION: 435

PRIOR APPLICATION DATA: including application PRIOR APPLICATION DATA: including application PRIOR APPLICATION DATA: described below:

APPLICATION NUMBER: 09/008,895

FILING DATE: January 19, 1993

APPLICATION NUMBER: 07/899,849

FILING DATE: December 7, 1992

ATTORNEY/AGENT INFORMATION:

NAME: Warburg Nichard J.

REFIERACE/DOCKET NUMBER: 208/149

TELECOMUNICATION INFORMATION:

TELECOMONICATION INFORMATION:

TELECOMONICATION INFORMATION:

TELECOMONICATION INFORMATION:

TELECAMONICATION INFORMATION:

TELECAMONICATION TORPORTON:

SEQUENCE CHARACTERISTICS:

"DANGTH: 15 DASE PAILE
                                                  Patent No. 5837542
GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dames McSwiggen
APPLICANT: Dames McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Renneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RIBOZYME APPLESION
TITLE OF INVENTION: MURRACELLULAR APPLESION
TITLE OF INVENTION: MOLECULE-1 (1-CAM-1)
NUMBER OF SEQUENCES: 2390
                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM COMPALIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620A
FILLING DATE: August 17, 1994
        US-08-292-620A-105/c Sequence 105, Application US/08292620A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-292-620A-393/c
) Sequence 393, Application US/08292620A
Patent No. 5837542
) GRABREAL INFORMATION:
APPLICANT: Susan Grimm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      308 GGGCTGCAACTCCAT 322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15 GGGCTGGAACCCCAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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Gaps

RESULT 475

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us-08-463_483A-34/c

j Sequence 34, Application US/08463483A

Patent No. 5849870

GENERAL INFORMATION:
APPLICANT: Warren, Gregory W
APPLICANT: Mullins, Martha A
APPLICANT: Oarr, Brian
APPLICANT: Carr, Brian
APPLICANT: Duck, Nicholas B
APPLICANT: Estruch, Juan J
APPLICANT: Bostichka, N. Kristy
APPLICANT: Bostichka, No. 5849870el Pesticidal Proteins and Strains
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: TURA-GEIGY Corporation
STREET: 7 Skyline Drive
CITY: Hawthorne
CITY: Hawthorne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 15;
                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/471,044
FILING DATE: 06-JUN-1995
CLASSIFICATION DATA:

PRICA APPLICATION DATA:

CLASSIFICATION DATA:

APPLICATION NUMBER: US/08/463,483
FILING DATE: 05-JUN-1995
PRICA APPLICATION DATA:

APPLICATION NUMBER: US/08/314,594
FILING DATE: 09-SEP-1994
PRICA PAPLICATION DATA:

APPLICATION NUMBER: US/08/218,018
FILING DATE: 23-MAR-1994
PRICA PAPLICATION DATA:

APPLICATION UNBER: US/08/037,057
FILING DATE: 25-MAR-1993
ATTORNEY/AGENT INFORMATION:

NAME: PROCE (BATY M.

NAME: PROCE (BATY M.

REGISTRATION NUMBER: US/08/037,057
FILING DATE: 25-MAR-1993
ATTORNEY/AGENT INFORMATION:

NAME: PROCE (BATY M.

SECURIOR OF SECULOR OF SECUL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%; Score 11.8; DB 1; Length 1
86.7%; Pred. No. 1.9e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /desc = "reverse primer used to make
pCIB5526"
                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          826 AIGCAGCIGAAGCIT 840
                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best_Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-471-044-34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               à
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APPLICANT: Koziel, Michael G
APPLICANT: Mullins, Martha A
APPLICANT: Mullins, Martha A
APPLICANT: Desai, Malini M
APPLICANT: Desai, Nalini M
APPLICANT: Kostichka, N. Kristy
APPLICANT: Betuch, Juan J
APPLICANT: A SEQUENCES: SO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      two
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                                                                                                                                                                                                                                                                          SIKERI: SULCE 4/700
CITY. LOS ANGELES
COUNTRY: LOS ANGELES
COUNTRY: LOS ANGELES
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" DISKETTE, 1.000
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620A
FILING DATE: AUGUST 17, 1994
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: described below:
APPLICATION NUMBER: 08/008,995
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
MARKEY/AGENT INFORMATION:

TITLE OF INVENTION: INTRACELLULAR ADHESION TITLE OF INVENTION: MOLECULE-1 (I-CAM-1) NUMBER OF SECURNCES: 2390 CORRESPONDENCE ADDRESS: ADONE & Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: BSILLUC...
TITLE OF INVENTION: No. 58400.
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: CIBA-GEIGY Corporation
""PEFT: 7 Skyline Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 477
US-08-471-044-34/c
; Sequence 34, Application US/08471044
; Patent No. 5440868
; GENERAL INFORMATION:
; APPLICANT: Warren, Gregory W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 656:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1260 CCAGGTTGAGGCCCT 1274
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS.
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ccasscreassrct 1
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Gaps

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TOPOLOGY: linear
MOLECULE TYPE: Genomic DNA
DESCRIPTION: gamma-crystallin gene exons 1 and 2
DESCRIPTION: (Accession # K03003) nucleotides 144 to 158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.9%; Score 11.8; DB 1; Length 15;
86.7%; Pred. No. 1.9e+02;
live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AUTHORS: Meakin, S O, Breitman, M L, Tsui, L C. TITLE: Structural and evolutionary TITLE: relationships among five members of the human JOURNAL: Molecular and Cellular Biology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: WANG, C. -G.
APPLICANT: WEBBURN, A. G.
TITLE OF INVENTION: GENETIC SEQUENCE ASSAY USING DNA
TITLE OF INVENTION: TRIPLE-STRAND FORMATION.
CORRESPONDENCE: 365
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 :FROM 1 TO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: PROFILE DIAGNOSTIC SCIENCES, INC., STREET: 510 EAST 73RD STREET,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 10021.
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44Mb storage
COMPUTER: IBM PC/XT/AT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: MS-DOS version 6.2 SOFTWARE: Wordperfect Version 5.1 CURRENT APPLICATION DATA:
FILLING DATE: 22 DEC 1993
CLASSIFICATION: 435
                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: (attorney) (212) 708-1880
TELEFAX: (attorney) (212) 246-8959
FFORMATION FOR SEQ ID NO: 61:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 07/968,436
FILING DATE: 29 OCT 1992
ATTORNEY/AGENT INFORMATION:
NAME: Handelman, Joseph H.
REGISTRATION NUMBER: 05/179
REFERENCE/DOCKET NUMBER: U9518-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HYPOTHETICAL: NO
ANTI-SENSE: NO
ORGOTHAL SOURCE:
ORGANISM: Home sapiens
POSITION IN GENOME:
CHROMOSOME/SEGMENT: chromosome 2
MAP POSITION: 2q33-q35
                                                                                                                                                                                                                                                 INFORMATION FOR ESO, ID NO: 61: 246-88
INFORMATION FOR ESO, ID NO: 61: SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: Nucleic Acid
STRANDENESS: double stranded
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 87, Application US/08173489C; Patent No. 5861244; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DATE: 1985
RELEVANT RESIDUES IN SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1142 CCTTTTTTTTTTT 1156
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Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PUBLICATION INFORMATION:
AUTHORS: Meakin, SO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5
1408-1414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NEW YORK
NEW YORK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-173-489C-61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-173-489C-87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JOURNAL: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-173-489C-61/c
| Sequence 61, Application US/08173489C
| Patent No. 2861244
| GENERAL INCORMATION:
| APPLICANT: WANG, C. -G.
| TITLE OF INVENTION: GENETIC SEQUENCE ASSAY USING DNA. TITLE OF INVENTION: TRIPLE-STRAND FORMATION.
| NUMBER OF SEQUENCES: 365
| CORRESPONDENCES: 365
| ADDRESSEE: PROFILE DIAGNOSTIC SCIENCES, INC., STREET: 510 EAST 73RD STREET,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "reverse primer used to make
DESCRIPTION: pCIB5526"
                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.30B
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/463,483A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRI: ...
ZIP: 10021.
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44Mb storage
COMPUTER: IBM PC/XT/AT
COMPUTER: IBM PC/XT/AT
SOFTWARE: Wordperfect Version 5.1
SOFTWARE: Wordperfect Version 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/173,489C
**TING DATE: 22 DEC 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGC 1695/CIP3
                                                                                                                                                                                                    PILLING DATE:
CLASSIFICATION: 530
PRICE APPLICATION: 530
PRICE ALGATION DATA:
APPLICATION NUMBER: US 08/314,594
PRICE APPLICATION DATA:
APPLICATION NUMBER: US 08/218,018
PRICE APPLICATION DATA:
APPLICATION NUMBER: US 08/218,018
PRICE APPLICATION DATA:
APPLICATION NUMBER: US 08/037,057
PILLING DATE: 25-MAR-1994
APPLICATION NUMBER: 25-943
ATTORNEY/AGENT INFORMATION:
NAME: SPRUILI, W. MALTARY
REGISTRATION NUMBER: 32,943
RETERENCE/DOCKET NUMBER: 32,943
REGISTRATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              826 ATGCAGCTGAAGCTT 840
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STATE: NEW YORK
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-463-483A-34
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86.7%; Pred. No. 1.9e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CGC1695/CIP3/DIV8 - SQLv4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    other nucleic acid
/desc = "reverse primer used to make
pCIB5526"
NO
                                              COMPUTER EACABABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30B
CAURENT APPLICATION DATA:
PILING DATE: 06-UUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 482
US-06-774-306A-201

Sequence 201, Application US/08774306A

Parent No. 5869523
GENERAL INFORMATION:
TITLE OF INVENTION: METHOD AND REAGENT FOR TITLE OF INVENTION: INHIBITING HEPALITIS C TITLE OF INVENTION: URUS REPLICATION
TITLE OF INVENTION: URUS REPLICATION
COMPARE OF SEQUENCES: 497
COMPASSORDENCE ADDRESS:
ADDRESSER: Lyon & Lyon
STREET: Suite 4700

STREET: Suite 4700
                                                                                                                                                                                                                                  CLASSIFICATION: 435
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/463,483
FILING DATE: 05-7UN-1995
FRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/314,594
FILING DATE: 09-SEP-1994
PRIOR APPLICATION NUMBER: US 08/218,018
FILING DATE: 23-MAR-1994
PRIOR APPLICATION NUMBER: US 08/037,057
FILING DATE: 25-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Meigs, J. Timothy
RESERRACE/DOCKET NUMBER: 38,241
REFERENCE/DOCKET NUMBER: 38,241
RELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 919-541-8689
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 826 ATGCAGCTGAAGCTT 840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15 AAGGAGCTGAAGCTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DESCRIPTION:
DESCRIPTION:
HYPOTHETICAL:
         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Farent No. 5866326
GENERAL INFORMATION:
APPLICANT: Warren, Gregory W
APPLICANT: Koziel, Michael G
APPLICANT: Mullins, Martha A
APPLICANT: Desai, Walini M
APPLICANT: Besai, Walini M
APPLICANT: Bestruch, Usan J
TILLE OF INVENTION: Method For Isolating Vegetative Insecticidal
TILLE OF INVENTION: Method For Isolating Vegetative Insecticidal
MUMBER OF SEQUENCES: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
0
                                                             NAME: Handelman, Joseph H.

REGISTRATION NUMBER: 26,179
REGISTRATION NUMBER: 19518-6
REIGENOMUNICATION INPORMATION:
TELEPONE: (attorney) (212) 708-1880
TELEPANE: (attorney) (212) 246-8959
TELEPAN: (attorney) (212) 246-8959
INFORMATION FOR SEQ ID NO: 87:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 Dase pairs
TYPE: Nucleic Acid STRANDEDNESS: double stranded
STRANDEDNESS: double stranded
MOLECULE TYPE: Genomic DNA
DESCRIPTION: M33647, J02994) nucleotides 4062 to 4076
HYPOTHETICAL: NO
HYPOTHETICAL: NO
MANT-CENORMERS: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.9%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 1.9e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                              HYPOTHETICAL: No
ANTI-SENSE: No
ORIGINAL SCURCE:
ORGANISM: Home sapiens
POSITION IN GENOME:
CHROWOSOWE/SEGMENT: chromosome 13
MAP POSITION: 1314.2
PUBLICATION INCRMATION:
AUTHORS: Friend, S H, Horowitz, J M, Gerber, M R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in retinoblastomas and mesenchymal tumors: Organization of the sequence and its encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RELEVANT RESIDUES IN SEQ ID NO: 87 : FROM 1 TO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Proceedings of the National Academy of Sciences, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: No. 5866326artis Corporation
STREET: 3054 Cornwallis Road
CITY: Research Triangle Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RA.
Deletions of a DNA sequence
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/968,436
FILING DATE: 29 OCT 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1202 CTTCACACCTCCCT 1216
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Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: NO. 58663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9059-9063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein
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JOURNAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AUTHORS:
TITLE: De
TITLE: ir
TITLE: O:
TITLE: PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VOLUME:
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Gaps

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                PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/218,018
FILING DATE: 23-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/037,057
FILING DATE: 25-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAMB: Medgs, J. Timothy
REGRENTATION NUMBER: CGC1695/CIP3/DIV4 - SQLV4
TELEPHONE 1919-541-8689
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 15;
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US-08-585-684B-775/c
; Sequence 775, Application US/08585684B
; Patent No. 5877021
; GENERAL INFORMATION:
; APPLICANT: Stinchcomb, Daniel T.
APPLICANT: McSwiggen, James
; TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
; TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
; NUMBER OF SEQUENCES: 2751
; CORRESPONDENCE ADDRESSE:
; ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
; STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "reverse primer used to make
DESCRIPTION: pCIB5526"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.9%; Score 11.8; DB 1; Best Local Similarity 86.7%; Pred. No. 1.9e+02; Matches 13; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STAKET: SULCE 4,000
CITY: LOS ANGELES
STATE: California
COUNTRY: U.S.A.
ZIP: 90071
ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastESE Varsion 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,684B
FILING DATE: January 16, 1996
APPLICATION NUMBER: 60/000,951
FILING DATE: July 7, 1995
ATTORNEY, AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELEDRONE: (213) 499-1600
TELEBRONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          826 ATGCAGCTGAAGCTT 840
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APPLICANT: Woziel, Michael G
APPLICANT: Wolfins, Martha A
APPLICANT: Mullins, Martha A
APPLICANT: Wolf Gordon J
APPLICANT: Osri, Brian
APPLICANT: Kostichka, N. Kristy
APPLICANT: Estruch, Juan J
APPLICANT: Estruch, Juan J
APPLICANT: No. 5872212el Pesticidal Proteins and Strains
NUMBER OF SEQUENCES: 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 11.8; DB 1; Length 15; 53.3%; Pred. No. 1.9e+02; tive 5; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.30B
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,566B
FILING DATE: 06-UIN-1995
CLASSIFICATION TATA:
APPLICATION NUMBER: US 08/463,483
FILING DATE: 05-UIN-1995
PRIOR APPLICATION NUMBER: US 08/463,483
FILING DATE: 05-UIN-1995
PRIOR APPLICATION DATE: US 08/314,594
FILING DATE: 09-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             No. 5872212artis Corporation
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/774,306A
FILING DATE: December 26, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/182,968
FILING DATE: Annuary 13, 1994
APPLICATION NUMBER: 07/882,888
FILING DATE: May 14, 1952
ATTORNEY,AGBNT INPORMATION:
NAME: WALDLY, RICHARTO:
NAME: WALDLY, RICHARTO:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 201:
SEQUENCE CHARACTERISTICS:
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-470-566B-34/c
; Sequence 34, Application US/08470566B
; Patent No. 5872212
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
BDIUM TYPE: Floppy disk
CMBUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 3054 Cornwallis Road
CITY: Research Triangle Park
STATE: NC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1190 TGAGTGTTGGACCTT 1204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 53.34
Matches 8, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-774-306A-201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: US
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US-08-585-684B-1365/c
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                                                                                                                                                                                               0.9%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                             US-08-589-684B-776/c
; Sequence 776, Application US/08585684B
; Patent No. 5877021
; GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Stinchcomb, Daniel T.
APPLICANT: McSwigen, James
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
TITLE OF SEQUENCES: 2751
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                           2; Indels
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0.9%; Score 11.8; DB 1;
Best Local Similarity 86.7%; Pred. No. 1.9e+02;
Matches 13; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: CALLIOLIDA

ZIP: 90071

ZIP: 90071

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 6torage
COMPUTER: IBM Compatible
OPERATURE SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/586,684B
FILING DATE: January 16, 1996
FILING DATE: January 16, 1996
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: WASDING: Richard
NAME: WASDING: Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
TELEPHONE: (213) 955-0440

INFORMATION FOR SEQ ID NO: 776:
SEQUENCE CHARRATERISTICS:
TUMCATH. 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                       1157 GGAAGTAAAGCAGCT 1171
                                                                                                                                                                                                                                                                                                                              15 GGAAGCAAAGCAGGT 1
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 15 base pairs TYPE: nucletc acid STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                    Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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US-08-585-684B-775
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Gaps
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Sequence 1365, Application US/08585684B
Patent No. 5877021
GENERAL INFORMATION:
APPLICANT: Strinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: MUDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES;
NUMBER OF SEQUENCES: 2751
CORRESPONDENCE: Lyon & Lyon
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APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
NUMBER OF SEQUENCES: 2751
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.9%; Score 11.8; DB 1;
86.7%; Pred. No. 1.9e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FractSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,684B
FILING DATE: January 16, 1996
FILING DATE: January 16, 1996
PRIOR APPLICATION NUMBER: 60/000,951
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1955-0440
TELEEX: 67-3510
TELEEX: 67-3510
TELEEX: 67-3510
TELECOMMUNICATION POR SEQ ID NO: 1365:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
ZIP: 90071
COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 1376, Application US/08585684B
; Patent No. 5877021
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                             ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1155 TTGGAAGTAAAGCAG 1169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid
EDNESS: single
                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: 1;
US-08-585-684B-1365
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APPLICANT: Pearson, Robert E.
APPLICANT: Dickson, Julie A.
APPLICANT: Mehrponyan, Majid
TITLE OF INVENTION: STRAND DISPLACEMENT AMPLIFICATION OF RNA
TITLE OF INVENTION: TARGETS
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSE: R. J. Rodrick, Becton Dickinson and Company
STREET: Becton Drive
CITY: Franklin Lakes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 1.9e+02; rive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: US
ZIP: 074.17
COMPUTER REABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/854,041
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match

0.9%; Score 11.8; DB 1;
Best Local Similarity 73.3%; Pred. No. 1.9e+02;
Matches 11; Conservative 2; Mismatches 2;
APPLICATION NUMBER: 60/000,951
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
TELEFAX: (213) 955-0440
TELEFAX: (213) 955-0410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      32,135
ER: P-3853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-854-041-4/c; Sequence 4, Application US/08854041; Patent No. 5916779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:

NAME: FUGIT, DOTHAR R.

REGISTRATION NUMBER: 32,135

REFERRACE/DOCKET NUMBER: P-

SEQUENCE CHARACTERISTICS:

LENGTH: 15 base pairs

TYPE: nucleic acid

STARNDEDNESS: single

'TOPOLOGY: linear

US-08-854-041-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           276 CAAAGAGGAAGCAGC 290
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Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                 ; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-585-6848-2270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE:
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0.9%; Score 11.8; DB 1; Length 15;
Best Local Similarity 33.3%; Pred. No. 1.9e+02;
Matches 5; Conservative 8; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2270, Application US/08585684B
Fatent No. 5877021
GENERAL INFORMATION:
APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Grinchcomb, Daniel T.
APPLICANT: MCSWiggen, James
TITLE OF INVENTION: MCSWiggen, James
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
NUMBER OF SEQUENCES: 2751
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: Saile 4700
                                                                                                                                                                             MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFWARE: FastESQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,684B
FILING DATE: January 16, 1996,
PRIOR APPLICATION NUMBER: 60/000,951
FILING DATE: July 7, 1995
APPLICATION NUMBER: 60/000,951
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: WARDLIGY, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION OF 1376:
SEQUENCE CHARACTERISTICS:
TEMATION FOR SEG ID NO: 1376:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
                                                                                                               ZIP: 90071
COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFWARE: FRESESQ Version 1.5
CURRENY APPLICATION DATA:
APPLICATION NUMBER: US/08/585,684B
FILING DATE: January 16, 1996
  633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1109 TAGITITICIGITIAN 1123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 UGGUUUCUGUCUAA 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4: 15 base pairs
nucleic acid
DEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 633 West Fifth
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
                        Suite 4700
                                            Los Angeles
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TYPE: nucleic aci
; STRANDEDNESS: sir
; TOPOLOGY: linear
US-08-585-684B-1376
                                                                                              U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 488
US-08-585-684B-2270
                                                                                           COUNTRY:
                                            CITY:
STATE:
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Sequence 24, Application US/08343998A

Sequence 24, Application US/08343998A

Patent No. 6020123

GENERAL INFORMATION:

APPLICANT: Sonigo, Pierre

APPLICANT: Brechot, Christian

APPLICANT: Brechot, Christian

APPLICANT: Courganard, Valerie

ITILE OF INVENTION: OLIGONUCLECTIDE SEQUENCES FOR THE AMPLIFICATION OF THE

ITILE OF INVENTION: OLIGONUCLEOTIDE SEQUENCES FOR THE HIV-2 AND SIV TYPE,

ITILE OF INVENTION: AND THEIR USES FOR IN VITRO DIAGNOSIS OF THE INFECTIONS

FILE REPERENCE: 2386.0066-01

CURRENT APPLICATION NUMBER: US/08/343,998A

CURRENT APPLICATION NUMBER: 07/820,600

SARLIER APPLICATION NUMBER: 07/820,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 15;
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                                                                                                                                                                                CONTRY: USA
ZIP: 10532
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30B
CURRENT APPLICATION DATA:
APPLICATION NOMBER: US/08/469,334
FILING DATE: 06-UNM-1995
CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDENDONO
TOPOLOGY: linear
MUBGCULE TYPE: other nucleic acid
DESCRIPTION: /desc = "reverse primer used to make
PROCESTIPTION: pCIBS526"
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0.9%; Score 11.8; DB 1;
Best Local Similarity 86.7%; Pred. No. 1.9e+02;
Matches 13; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/463,483
FILING DATE: 09-SEP-1994
FILING DATE: 23-MAR-1994
FILING DATE: 23-MAR-1994
FILING DATE: 23-MAR-1994
FILING DATE: 25-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spruill, W. MUTTEY
REGISTRATION NUMBER: 32,943
REGISTRATION NUMBER: 32,943
REFERENCE/DOCKET NUMBER: CGC 1695/CIP3
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                          CORRESPONDENCE ADDRESS:
ADDRESSEE: CIBA-CBIGY Corporation
STREET: 7 Skyline Drive
CITY: Hawthorne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         826 ATGCAGCTGAAGCTT 840
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: sing
NUMBER OF SEQUENCES:
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US-08-343-998-24
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                                                                                                                           CITY:
STATE:
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                                                                                             Patent No. 597(799)

APPLICANT Allibert, Patrice A. APPLICANT Machand Bernard F. APPLICANT Machand Bernard F. APPLICANT Machandrand Bernard F. APPLICANT Machandrand Bernard F. APPLICANT Machandrand Bernard F. APPLICANT Machandrand Bernard F. APPLICANT Tiercy, Jean-Marit F. APPLICANT Tiercy, Jean-Marit F. APPLICANT Tiercy, Jean-Marit F. APPLICANT Tiercy, Jean-Marit F. BEREORMED, AND TYPING METHOD USING SAID PROBES CITY. Alexandria BEREORMED, AND TYPING METHOD USING SAID FROMESSER: OLIFF & BEREORMED, AND TYPING METHOD USING SAID FROMESSER: OLIFF & BEREORMED, AND TYPING METHOD USING SOURTING STREET: P.O. Box 1928

COMPUTER: READABLE FORM: MEDIA TRANSPARIE FORM: MEDIA TYPING SYSTEM: PC-DOS/MS-DOS SOURTING SYSTEM: PC-DOS/MS-DOS SOURTING SYSTEM: PC-DOS/MS-DOS SOURTING SYSTEM: PC-DOS/MS-DOS SOURTING DATE: T-UTW-1995

CLASSIFICATION NUMBER: US 08/030,143

FILING DATE: 11-MAR-1993

CLASSIFICATION NUMBER: US 08/030,143

FILING DATE: 11-MAR-1993

CLASSIFICATION NUMBER: US 08/030,143

FILING DATE: 11-MAR-1993

CLASSIFICATION NUMBER: 30,028

REGISTRATION NUMBER: 30,028
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Duck, Nicholas B
APPLICANT: Estruch, Juan J
TITLE OF INVENTION: No. 5990383el Pesticidal Proteins and Strains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NECOLA 7-34/C

1 Sequence 34, Application US/08469334

2 Sequence 34, Application US/08469334

3 Fatent No. 5990383

3 APPLICANT: Warren, Gregory W
APPLICANT: Koziel, Michael G
APPLICANT: Mullins, Martha A
APPLICANT: Mye, Gordon J
APPLICANT: Nye, Gordon J
APPLICANT: Nein A
APPLICANT: Nein A
APPLICANT: Kostichka, N. Kristy
APPLICANT: Kostichka, N. Kristy
APPLICANT: Desai, Nalini M
APPLICANT: Desai, Nalini M
APPLICANT: Desai, Nalini M
APPLICANT: Notholas B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              782 TCTCCACCAGTGCCC 796
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 703-836-640
TELEFAX: 703-836-2787
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
                                                                                   JS-08-485-133-7/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-485-133-7
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Gaps . 0 ö

Gaps

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Sequence 43, Application US/08832021
| Patent No. 604598
| Patent No. 604598
| GENERAL INFORMATION:
| APPLICANT: Combates, N. |
| APPLICANT: Pardinas, J. |
| APPLICANT: Pardinas, J. |
| APPLICANT: Proutho, S. |
| APPLICANT: PROUTHOR NUMBER: US/08/832,021 |
| CURRENT APPLICATION NUMBER: US/08/832,021 |
| NUMBER OF SEQ ID NOS: 64 |
| SOFTWARE: Patentin Ver. 2.0 |
| SOFTWARE: Patentin Ver. 2.0 |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 41, Application US/08632021
Patent No. 604599
GENERAL INFORMATION:
APPLICANT: Combates, N.
APPLICANT: Pardinas, J.
APPLICANT: Promity, S.
APPLICANT: PROMITY S.
APPLICANT S.
APPLICANT: PROMITY S.
APPLICANT S.
APPLICANT S.
APPLICANT S.
APPLICANT S.
APPLICANT S.
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US-08-832-021-43
                                    ; OTHER INFORMATION: Description of Artificial Sequence: primer US-08-832-021-37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: primer
                                                                                                                                          Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%; Score 11.8; DB 1;
86.7%; Pred. No. 1.9e+02;
tive 0; Mismatches 2;
                                                                                                                                          0.9%; Score 11.8; DB 1;
ilarity 86.7%; Pred. No. 1.9e+02;
Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 11.8; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/08/832,021
CURRENT FILING DATE: 1997-04-02
NUMBER OF SEQ ID NOS: 64
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 41
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1145 TTTTTTTTTTGGA 1159
                                                                                                                                                                                                                                                                                    TGGA 1159
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                     rrrrrrrrrrrsca 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.9
Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13; Conservative
                                                                                                                                                                                                                                                                                    1145 TTTTTTCTTT
                                                                                                                           Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
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US-08-832-021-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-832-021-41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-832-021-41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 43
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Sequence 25, Application US/08832021

Patent No. 6045998

GENERAL INFORMATION:
APPLICANT: Pardinco, S.
APPLICANTON: IMPROVED TECHNIQUE FOR DIFFERENTIAL DISPLAY
CURRENT FILING DATE: 1997-04-02

NUMBER OF SEQ ID NOS: 64
SOUTHARE: Patentin Ver. 2.0
SEQ ID NO 25
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .
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APPLICANT: Combates, N.

APPLICANT: Pardines, J.

APPLICANT: Prouty, S.

CURRENT SPERICATION NUMBER: US/08/832,021

CURRENT APPLICATION NUMBER: US/08/832,021

CURRENT FILING DATE: 1997-04-02

NUMBER OF SEQ ID NOS: 64

SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
) CTHER INFORMATION: Description of Artificial Sequence: primer US-08-832-021-25
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                                                                                                                                                                                                                                                                                                                                                                                         Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                      Query Match

0.9%; Score 11.8; DB 1;
Best Local Similarity 86.7%; Pred. No. 1.9e+02;
Matches 13; Conservative 0; Mismatches 2;
EARLIER APPLICATION NUMBER: PCT/FR90/00394
EARLIER FILING DATE: 1990-06-05
NUMBER OF SEQ ID NOS: 25
SEQ TWARE: PatentIn Ver. 2.0
SEQ ID NO 24
LENGTH: 15
                                                                                                                                                                                                                                            ORGANISM: Simian immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 37, Application US/08832021 Patent No. 6045998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                               TYPE: DNA
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1145 TTTTTTTTTGGA 1159
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Warren
US-08-832-021-61
                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Squence 47, Application US/08832021
Squence 47, Application US/08832021
Fatent No. 6045998
GENERAL INFORMATION:
APPLICANT: Combates, N.
APPLICANT: Pardinas, J.
APPLICANT: Pardinas, J.
APPLICANT: Prouty, S.
APPLICANT: Prouty, S.
TITLE OF INVENTION: IMPROVED TECHNIQUE FOR DIFFERENTIAL DISPLAY
FILE REPERENCE: JBD-382
CURRENT APPLICATION NUMBER: US/08/832,021
CURRENT FILIG DATE: 1997-04-02
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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0
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APPLICANT: Prouty, S.
APPLICANT: Prouty, S.
APPLICANT: Prouty, S.
APPLICANT: Stemm, K.
TITLE OF INVENTION: IMPROVED TECHNIQUE FOR DIFFERENTIAL DISPLAY
FILE REFERENCE: JBP-382
CURRENT APPLICANTON NUMBER: US/08/832,021
CURRENT FILING DATE: 1997-04-02
NUMBER OF SEQ ID NOS: 64
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Description of Artificial Sequence: primer US-08-832-021-45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 11.8; DB 1; Length 15; larity 86.7%; Pred. No. 1.9e+02; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.9%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 1.9e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                     US-08-832-021-45

Sequence 45, Application US/08832021

Patent No. 604598

GENERAL INFORMATION:

APPLICANT: Combates, N.

APPLICANT: Pardinas, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1145 TTTTTTTTTGGA 1159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1144 TTTTTTTTTTGG 1158
               1144 TTTTTTTTTGG 1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence
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Matches 13; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13;
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                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 45
LENGTH: 15
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LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
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Matches
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RESULT 499

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CENTRAL INFORMATION:
APPLICANT: PROLEY.
APPLICANT: PARTIANS.
TITLE OF INVESTION: 198-000-108-300-201
TITLE OF INVESTION: 198-000-108-300-201
TITLE OF INVESTION: 198-000-108-300-201
TITLE OF INVESTION: 198-000-108-300-201
TITLE OF INVESTION: 198-000-108-300-301
TITLE OF INVESTION: 198-000-108-300-301
TITLE OF INVESTION: 198-000-108-301
TITLE DATE: 198-000-108-301
TITLE
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0.9%; Score 11.8; DB 1; Length 15;
Best Local Similarity 53.3%; Pred. No. 1.9e+02;
Matches 8; Conservative 5; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Susan Grimm
APPLICANT: Susan Grimm
APPLICANT: Susan Grimm
APPLICANT: Sean Sullivan
APPLICANT: MUNENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
CORRESPONDENCES: 330
CORRESPONDENCES: 330
ADDRESSEE: Lyon & Lyon
STREET: Sulte 470
CITY: Los Angeles
STRRET: Galifornia
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER: IBM Compatible
OFFRATION SYSTEM: IBM P.C. DOS 5.0
SOFFWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE: January 19, 1993
APPLICATION NUMBER: US/08,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: January 19, 1993
APPLICATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
FELENCOMMULCATION INGRER: 208/149
FELENCOMMULCATION INGRER: 208/149
FELENCOMMULCATION INGRER: 208/149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 74, Application US/09071845
Patent No. 6132967
**LEFAX: (213) 489-1600
| TELEFAX: (213) 955-0440
| TELEX: 67-3510
| INFORMATION FOR SEQ ID NO: 201:
| SEQUENCE CHARACTERISTICS:
| LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (213) 489-1600
TELERX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 74:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                               1190 TGAGTGTTGGACCTT 1204
                                                                                                                                                                                                                                                                                                                                                                                                                              1 UGUGUGUUGGACCGU 15
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nucleic acid
                                                                                                                                                                     TYPE: nucleic acid

STRANDEDNESS: single

// TOPOLOGY:

US-09-064-156A-201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                              | DESCRIPTION: Other nucleic acid | DESCRIPTION: | Jesc = "reverse primer used to make | DESCRIPTION: pCIB5526" | HYPOTHETICAL: NO | US-09-300-529-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 201, Application US/09064156A
Fatent No. 6132966
Fatent No. 613296
Fatent No. 613296

GENERAL INPORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: VIRUS REPLICATION
TITLE OF INVENTION: VIRUS REPLICATION
NUMBER OF SEQUENCES: 499
CORRESPONDENCE: 499
CORRESPONDENCE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 470
CITY: Los Angeles
STATE: California
COUWTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.998
MEDIUM TYPE: ACORAGE
COMPUTER: BM Compatible
COMPUTER: BM Compatible
COMPUTER: BM Compatible
COMPUTER: APPLICATION DATA:
TILNG DATE: APPLICATION DATA:
APPLICATION NUMBER: 08/774,306
FILING DATE: December 26, 1996
APPLICATION NUMBER: 08/182,968
FILING DATE: January 13, 1994
APPLICATION NUMBER: 07/882,888
FILING DATE: MAN, 14,1007
                               PADLICATION NUMBER: US 08/218,018
FILING DATE: 23-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/037,057
FILING DATE: 25-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Meigs, J. Timothy.
REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: S-19506L
TELEPHONE: 919-541-8689
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: MUCLAIC COLD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 234/083
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        826 ATGCAGCTGAAGCTT 840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15 AAGGAGCTGAAGCTT 1
                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-064-156A-201
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                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Susan Grimm
APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME REATMENT OF
TITLE OF INVENTION: RIBOZYME ADHESION
TITLE OF INVENTION: NUTRACELLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SUSUBNESS: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Suite 4700
CITY: Los Angeles
STREET: Galifornia
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TILING DATE:

CLASSIFTCATION

CLASSIFTCATION

PRIOR APPLICATION NUMBER: US/08/292,620

FILING DATE: August 17, 1994

APPLICATION NUMBER: 08/008,895

FILING DATE: January 19, 1993

APPLICATION NUMBER: 07/989,849

FILING DATE: December 7, 1992

ATTORNEY/AGENT INFORMATION:

NAME: Warburg, Richard J.

REGISTRATION NUMBER: 2.08/149

FELECOMMUNICATION INFORMATION:

TELEFAX: (213) 489-1600

TELEFAX: (213) 955-0440

TELEFAX: (7.131) 955-0440

TELEFAX: (7.13
                                                                                                                                                                                                                                                US-09-071-845-393/c
; Sequence 393, Application US/09071845
; Patent No. 6132967
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1260 CCAGGTTGAGGCCCT 1274
                                                      308 GGGCTGCAACTCCAT 322
                                                                                                                 15 GGGCTGGAACCCCAT 1
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Best Local Similarity 86.7,
Matches 13; Conservative
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                                                                                                                                                                                                                 Gaps
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                                                                                                                                                Length 15;
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                                                                                                                                                                                                          2; Indels
                                                                                                                               Query Match

0.9%; Score 11.8; DB 1;
Best Local Similarity 73.3%; Pred. No. 1.9e+02;
Matches 11; Conservative 2; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Sean Grimm
APPLICANT: Damplication US/09071845
Fatent No. 6132967;
GENERALI INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dames McSwiggen
APPLICANT: James McSwiggen
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 MD
MEDIUM TYPE: 3.5" Diskette, 1.44 MD
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
CLASSIPICATION:
PRICR APPLICATION:
PRICR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: Dacember 7, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
APPLICATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELECOMUNICATION INFORMATION:
TELECHOMNICATION INFORMATION:
TELECHOMNICATION: 1015:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                 908 CCCTGGTCCTAAAGG 922
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STRANDEDNESS: sir
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Best Local Similarity
Matches 13; Conserv
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                                     ; TOPOLOGY:
US-09-071-845-74
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Length 15;
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       METHOD AND REAGENT FOR THE INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: Jarvis, Thale
APPLICANT: Jarvis, Thale
APPLICANT: Journal James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
NUMBER OF SEQUENCES: 2751
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.9%; Score 11.8; DB 1; Best Local Similarity 86.7%; Pred. No. 1.9e+02; Matches 13; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                          COUNTRY: U.S.A.
ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/038,073
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MEDIUM TYPE: storage
COMPUTER: IBM Compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 776, Application US/09038073; Patent No. 6194150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
PRICE APPLICATION DATA:
APPLICATION NUMBER: 08/585,684
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFRENCE/POCKET NUMBER: 32,327
REFRENCE/POCKET NUMBER: 218/(
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 955-0440
TELEPHONE: (713) 955-0440
TELERX: (713) 955-0440
TELERY: (713) 955-0440
TELERY: (713) 955-0440
TELERY: (713) 955-0440
                                                                                                                               ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1157 GGAAGTAAAGCAGCT 1171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15 GGAAGCAAAGCAGGT 1
                                                                                 2751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-09-038-073-775
                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Ly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
       TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                    CITY: Los Angeles
STATE: California
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STATE: California
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               Sequence 656, Application US/09071845
Fatent No. 6132967
GENERAL INFORMATION:
APPLICANT: Busan Grimm
APPLICANT: James McSwiggen
APPLICANT: Gens Sullivan
APPLICANT: Sean Sullivan
APPLICANT: Fenneth G. Draper
TITLE OF INVENTION: RIBGARED TO LEVELS OF
TITLE OF INVENTION: INTERACEL OF
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF SURVENTION: MOLECULE-1 (1-CAM-1)
MIMMED OF SURVENTION: MOLECULE-1 (1-CAM-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
BAPLICATION NUMBER: US/09/071,845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TILING DATE:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
RECISTRATION NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEFANE, (213) 955-0440
TELEFAX: 67-3510
INFORMATION FOR SEQ ID NO: 656:
                                                                                                                                                                                                              TITLE OF INVENTION: RIBGZERE TREATITLE OF INVENTION: DISEASES OR CONTINUE OF INVENTION: RELATED TO LETHILE OF INVENTION: RELATED TO LETHILE OF INVENTION: INTRACELLULAR TITLE OF INVENTION: MOLECULE-1 (I. NUMBER OF SEQUENCES: 2390 CORRESPONDENCE ADDRESS: ADDRESS: ADDRESS: STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 775, Application US/09038073
Patent No. 6194150
GENERAL INFORMATION:
APPLICANT: StinchComb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1260 CCAGGTTGAGGCCCT 1274
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STRANDEDNESS: single
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Matches 13; Conserv
US-09-071-845-656/c
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US-09-038-073-775/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-071-845-656
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                                                                                                                                                                                                                                                                                                                                  0.9%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 1.9e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%; Score 11.8; DB 1; Length 15; 33.3%; Pred. No. 1.9e+02; very 8; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
NUMBER OF SEQUENCES: 2751
CORRESPONDENCES: 2751
CORRESPONDENCES: ADDRESS: ADDRESSE: Lyon & Lyon
STREET: Suite 4700
CITY: Los Angeles
CITY: Los Angeles
STATE: Callfornia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
SOFTWARE: FABSERQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/038,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICALL...

RILLING DATE:
ATTORNEY AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELERA: (713) 955-0440
TELERA: 67-3510
INFORMATION FOR SEQ ID NO: 1376:
SEQUENCE CHARACTERISTICS:
"WATH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1376, Application US/09038073 Patent No. 6194150 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/585,684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1155 TIGGAAGTAAAGCAG 1169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1109 TAGTTTTCTGTTTAA 1123
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                                      SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLLOGY: linear
US-09-038-073-1365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15 TTGGAAGTACAGCTG 1
                                                                                                                                                                                                                                                                                                                                  Query Match 0.9
Best Local Similarity 86.7
Matches 13; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
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        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
Matches 5; Conserva
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0.9%; Score 11.8; DB 1; Length 15;
Best Local Similarity 86.7%; Pred. No. 1.9e+02;
Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Jarvis, Thale
APPLICANT: Jarvis, Thale
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
OURRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: Suite 4700
CITY: Los Angeles
STREET: Suite 4700
CITY: Los Angeles
STREET: Suite 8001
COUNTRY: US.A.
ED001
COMPTTER READABLE FORM:
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastESQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION ADDRESS: ADDRESS: ADDRESS: APPLICATION DATA:
APPLICATION ADDRESS: ADD
                                                                                                          APPLICATION NUMBER: US/09/038,073
FILING DATE:
APPLICATION NUMBER: US/09/038,073
APPLICATION NUMBER: 08/585,684
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REPERSORAPIONER 32,327
REPERSORAPIONER 32,327
TELEPHONE: (213) 499-1600
TELEPHONE: (213) 499-1600
TELEX: 67-5510
TELE
OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-038-073-1365/c
; Sequence 1365, Application US/09038073
; Patent No. 6194150
; GENERAL INFORMATION:
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APPLICATION NUMBER: 08/585,684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard
REGISTRATION UNDRER: 32,327
REFERENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
TELEFONE: (213) 489-1600
TELEFAX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1157 GGAAGTAAAGCAGCT 1171
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US-09-038-073-776
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Patent No. 6333152

j GRNERAL INFORMATION:
APPLICANT: Kinzler, Kenneth
APPLICANT: Abang, Lin
APPLICANT: Zhang, Lin
APPLICANT: Zhang, Lin
APPLICANT: Zhang, Lin
APPLICANT: Abang, Lin
APPLICANT: Abou, Wei

TITLE OF INVENTION: Gencer Cells
TITLE OF INVENTION: Gencer Cells
TITLE REFERENCE: 01107.74664

CURRENT APPLICATION NUMBER: US/09/081,646

CURRENT APPLICATION NUMBER: 60/047,352

EARLIER RILING DATE: 1997-05-21

NUMBER OF SEQ ID NOS: 871

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 513

LENGTH: 15
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                                                                                                     Length 15;
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0.9%; Score 11.8; DB 1;
Best Local Similarity 86.7%; Pred. No. 1.9e+02;
Matches 13; Conservative 0; Mismatches 2;
                                                                                                       0.9%; Score 11.8; DB 1;
86.7%; Pred. No. 1.9e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                             | Facell NIN. 042443.
| Facell NIN. 042443.
| APPLICANT: KOPETZKİ, ETHAYĞ
| APPLICANT: KOPETZKİ, ETHAYĞ
| APPLICANT: SCHANT: Christian
| TITLE OF INVENTION: New Host-Vector System
| FILE REFERENCE: CD20315
| CURRENT APPLICATION NUMBER: US/09/344,888A
| CURRENT APPLICATION NUMBER: EP98113156.8
| PRIOR FILING DATE: 1998-06-25
| PRIOR PILING DATE: 1998-07-15
| PRIOR PILING DATE: 1998-10-09
| NUMBER OF SEQ ID NOS: 24
| SOFTWARE: PATENTIN VET. 2.0
| SEQ ID NO 9
                                                                                                                                                                                                                                                                                  US-09-344-888A-9/c
; Sequence 9, Application US/09344888A
; Patent No. 6291245
                                                                                                                                                                             719 CCCAGCAGCAGGGG 733
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ORGANISM: Artificial Sequence
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Best Local Similarity 86.73
Matches 13; Conservative
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ORGANISM: Homo sapiens
                               TYPE: RNA
CRGANISM: E. coli
US-09-275-850-19
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SEQ ID NO 19
LENGTH: 15
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                                       Sequence 19, Application US/09275850A
Patent No. 6261774
GENERAL INFORMATION:
APPLICANT: Pagratis, Nikos
APPLICANT: Gold, Larry
APPLICANT: Statland, Timur
APPLICANT: Javornik, Brenda
ITILE OF INVENTION: Truncation SELEX Method
FILE REFERENCE: NEX 79
CURRENT FILING DAIE: 1999-03-24
NUMBER OF SEQ ID NOS: 351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Warburg Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/585,684
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 73.3
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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             RESULT 510
US-09-038-073-2270
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US-09-275-850-19
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PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: 1BM Compatible
OPERATING SYSTEM: 1BM MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/06331A
                                                                                                                                                                                                                  Pharmaceuticals, Inc.
METHOD AND REAGENT FOR
TREATMENT OF FIBROSIS P
FIBROUS TISSUE DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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June 2, 1994
                                                                                                                                                                                                             APPLICANT: Ribozyme Pharmaceuticals, TITLE OF INVENTION: METHOD AND REAGEN TITLE OF INVENTION: TREATMENT OF FIBH TITLE OF INVENTION: FIBROUS TISSUE DI NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                  PCT-US94-06331A-9/c; Sequence 9, Application PC/TUS9406331A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                E: Lyon & Lyon
611 West Sixth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ٥.
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20,
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: (7-3510
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        688 GGGAGCCAGCGCCC 702
                           334 CCTGGTGATAGTCAC 348
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Best Local Similarity 86.7
Matches 13; Conservative
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CITY: Los Angeles
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nucleic acid
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PCT-US94-06331A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: Juctassification:
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GENERAL INFORMATION:
APPLICANT: Kinzler, Kenneth
APPLICANT: Kinzler, Kenneth
APPLICANT: Lin
APPLICANT: Zhou, Wei
TITLE OF INVENTION: Gene Expression Profiles in No. 633152mal and
TITLE OF INVENTION: Gancer Cells
TITLE OF INVENTION: Gancer Cells
FILE REFERENCE: 01107.74664
CURRENT APPLICATION NUMBER: 05/091,646
CURRENT FILING DATE: 1998-05-20
EARLIER FILING DATE: 1998-05-21
NUMBER OF SEQ ID NOS: 871
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 616
LENGTH: 15
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                                             Gaps
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Fatent No. 6472586

GENERAL INFORMATION:

APPLICANT: Maliga, Pal

APPLICANT: Hasdwicz, Peter T.

APPLICANT: Hasdwicz, Peter T.

APPLICANT: Hasdwicz, Peter T.

APPLICANT: Hasdwicz, Peter T.

APPLICANT: Rutgers, The State University of New Jersey

TITLE OF INVENTION: Plastids of Higher Plants

FILE NEFERENCE: Rut-95-08031

CURRENT APPLICATION NUMBER: US/09/011,336

CURRENT APPLICATION NUMBER: PCT/US96/12671

PRIOR FILING DATE: 1998-08-01

PRIOR FILING DATE: 1995-08-01

PRIOR FILING DATE: 1995-08-10

NUMBER OF SEQ ID NOS: 70

SOFTWARE FERENCE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 1.9e+02; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 15;
Query Match 0.9%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 1.9e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3; 1.9e+02; 1.9e+02; 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.9%; Score 11.8; D'
86.7%; Pred. No. 1.9e
7ative 0; Mismatches
                                                                                                                                                                                                                                    Sequence 616, Application US/09081646 Patent No. 6333152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1289 CAGTIGCTCAGCCTG 1303
                                                                                   1240 CIGGACGIGGCCAIG 1254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: PCR Primer US-09-011-336-23
                                                                                                                         15 crccáceredecare 1
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Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                                                                                                                                                                                                               -09-081-646-616/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-011-336-23/c
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LENGTH: 15
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0.9%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                              Patent No. 5182195

APPLICANT: NAKHAMA, KAZUO; KAISHO, YOSHIHIKO; YOSHIMURA, K;
TITLE OF INVENTION: METHOD FOR INCREASING USING PROTEASE
DEFICIENT YEASTS

UNMBER OF SEQUENCES: 71

CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/269,140

FILING DATE: 09-NOV-1988
                                                Indels
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COMPUTER READABLE PORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage COMPUTER: 1BM Compatible OPERATURG SYSTEM: 1BM MS-DOS (Version 5.0) COMPRENT MORTHER: Wordberfect (Version 5.1) CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/23,030
FILING DATE:
CLASSIFICATION: 536
PRIOR PEPLICATION DATA:
APPLICATION NUMBER: US/08/008,910
FILING DATE:
APPLICATION NUMBER: US/08/23,030
FILING DATE:
APPLICATION NUMBER: 32,327
REFERENCE/POCKET NUMBER: 32,327
REFERENCE/POCKET NUMBER: 197/240
TELECOMMUNICATION INFORMATION:
TELECHONE: (213) 489-1600
TELECOMMUNICATION INFORMATION:
TELECHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match

0.9%; Score 11.8; DB 1;
Best Local Similarity 73.3%; Pred. No. 2.38+02;
Matches 11; Conservative 2; Mismatches 2;
METHOD AND REAGENT FOR
TREATMENT OF PROMYELOCYTIC
LEUKEMIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 780, Application US/08291932A

Sequence 780, Application US/08291932A

Patent No. 5658780

GENERAL INFORMATION:
APPLICANT: Stinchcomb, Dan T.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: NF-XB

TITLE OF INVENTION: NF-XB

TITLE OF INVENTION: NF-XB

NUMBER OF SEQUENCES: 830

CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700

CITY: Los Angeles
STREET: Galifornia
COUNTRY: U.S.A.
ZIP: 90071-206

COMPUTER REALABLE FORM:
MEDIUM TYPE: Storage

COMPUTER: IBM Compatible
  TITLE OF INVENTION: METHOD AND REZ
TITLE OF INVENTION: TERRATMENT OF I
TITLE OF INVENTION: LEUKEMIA
NUMBER OF SEQUENCES: 62
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 611 West Sixth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1053 CAGCCCTGGCCTTCC 1067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (213) 955-044
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TYPE: nucleic acid
; STRANDEDNESS: singl
; TOPOLOGY: linear
US-08-233-030-52
                                                                                                                                                                          CITY: Los Angeles
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 520
US-08-291-932A-780
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                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Weiss, Arthur
APPLICANT: Weiss, Arthur
APPLICANT: Weiss, Arthur
APPLICANT: Weiss, Arthur
APPLICANT: Irving, Brian A.
APPLICANT: Zeabo, Kristina
TITLE OF INVENTION: Chimeric Chains for Receptor
TITLE OF INVENTION: Associated Signal Transduction Pathways
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton &
ADDRESSEE: Flehr, Hobbach
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
                                                                                                         0
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                                                 Query Match

0.9%; Score 11.8; DB 1; Length 15;
Best Local Similarity 86.7%; Pred. No. 1.9e+02;
Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTY: USA

ZIP: 9411-4187

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PATEMIT Release #1.0, Version #1.25
SOFTWARE: PATEMIT Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/988,194A
FILING DATE: December 9, 1992
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: ROALAND, BERTRAIL 20015
REGISTRATION NUMBER: 20015
REGISTRATION NUMBER: 20015
REGISTRATION NUMBER: 20015
REGISTRATION NUMBER: 20015
REGISTRATION NUMBER: 20015
REGISTRATION NUMBER: 20015
REGISTRATION NUMBER: 20015
REGISTRATION NUMBER: 20015
REGISTRATION NUMBER: 20015
REGISTRATION SEQ ID NO: 16: SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: MUCLeic acid
TYPE: MUCLeic acid
TYPE: MUCLeic acid
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US-08-233-030-52
Sequence 52, Application US/08233030
FATERIT NO. 5639655
GENERAL INFORMATION:
APPLICANT: James D. Thompson
APPLICANT: Kenneth G. Draper
                                                                                                                                                                                                                                                                                                   US-07-988-194A-16/c
; Sequence 16, Application US/07988194A
Patent No. 5359046
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1239 GCTGGACGTGGCCAT 1253
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       S
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                                                                                              13;
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STATE: Cl
COUNTRY:
                                                                                                                                                                                                                                                                           RESULT 518
  5182195-24
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Gaps

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Length 16; Indels us09904568-3.rni

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APPLICANT: FINER, MITCHELL H.
APPLICANT: FINER, MITCHELL H.
APPLICANT: ROBERTS, MARGO R.
APPLICANT: DULL, THOMAS J.
APPLICANT: ZSEBO, KRISZINA M.
APPLICANT: ZSEBO, KRISZINA M.
APPLICANT: QIN, LU
TITLE OF INVENTION: WETHOD FOR PRODUCTION OF HIGH TITER
TITLE OF INVENTION: OF MANMALIAN CELLS
NUMBER OF SEQUENCES: 32
CORRESPONDRACE ADDRESS:
ADDRESSEE: CELL GENESYS, INC.
STREET: 322 LAKESIDE DRIVE
CITY: ROSIER CITY
STATE: CALLFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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MEDIUM TYPE: FIOPPY disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/258,152

FILING DATE: 10-JUN-1994

CLASSIFICATION OATA:

APPLICATION NUMBER: US 08/076,299

FILING DATE: 11-JUN-1993

ATTORNEY/AGENT INFORMATION:

REGISTRATION NUMBER: 34,647

RELEPHONE: 415-358-9600 X131

TELEFRAK: 415-358-9600 X131

TELEFRAK: A15-359-352
FILING DATE: May 18, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/157
TELECOMMUNICATION INFORMATION:
TELEPAK: (213) 489-1600
TELEPAK: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 814:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
TYPE: MUCLEIC SEG ID
STENGTH: 16 base pairs
TYPE: MUCLEIC SEG ID
STENGTH: 16 base pairs
TYPE: MUCLEIC SEG ID
STENGTH: 16 base pairs
TYPE: MUCLEIC SEG ID
STENGTH: 16 base pairs
TYPE: MUCLEIC SEG ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18, Application US/08258152 Patent No. 5686279
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LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 415-349-7392
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
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US-08-291-932A-814
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 522
US-08-258-152-18/c
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                                                                      PILING DATE: August 1.,

CLASSIFICATION 514

PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994

APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994

ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327

REFERENCE/DOCKET NUMBER: 208/157

TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1800

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Sequence 814, Application US/08291932A

Patent No. 5688780

GENERAL INFORMATION:

APPLICANT: Stinchcomb, Dan T.

APPLICANT: Draper, Kenneth G.

APPLICANT: McSwiggen, James

TITLE OF INVENTION: RIBOZYMB TREATMENT OF

TITLE OF INVENTION: RIBOZYMB TREATMENT OF

TITLE OF INVENTION: RIELATED TO LEVELS OF

TITLE OF INVENTION: RELATED TO LEVELS OF

TITLE OF INVENTION: NF-KB

NUMBER OF SEQUENCES: 830

CORRESPONDENCE ADDRESS:

ADDRESSER: Lyon & Lyon

STREET: 633 West Fifth Street

STREET: Suite 4700

CITY: Los Angeles

COUNTRY: U.S.A.

ZIP: A.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 1BM COMPATIBLE
COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
COMPUTER: Word Perfect 5.1
CURRENT APPLICATION DATA:
FILING DATE: August 15, 1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
PRIOR APPLICATION DATA: described below:
APPLICATION NUMBER: 08/245,466
                               SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/291,932A
FILING DATE: August 15, 1994
           OPERATING SYSTEM: IBM P.C. DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPAL...
TELEPAK: (21s)
TELERA: 67-3510
INFORMATION FOR SEQ ID NO: 78
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TVPE: mucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           480 GGACTGCCGAGACGG 494
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12; Conservative
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Matches 12; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
US-08-291-932A-780
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US-08-291-932A-814
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GENERAL INFORMATION:
APPLICANT: Reed, John
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                                                                                                                                                                                                               COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ANTI-SENSE:
US-08-465-485A-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             à
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                                                                                                                                    Gaps
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                                                                                                                                  .
                                                                                     Query Match 0.9%; Score 11.8; DB 1; Length 16; Best Local Similarity 86.7%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Jun KONDO
APPLICANT: Atsuco KOHARA
APPLICANT: Atsuco KOHARA
APPLICANT: Ei YAMADA
TITLE OF INVENTION: HUMAN CHONDROMODULIN-I PROTEIN
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA.
APPLICATION NUMBER: US/08/241,465B FILING DATE: MAY 11, 1994 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.9%; Score 11.8; DB 1;
Best Local Similarity 86.7%; Pred. No. 2.3e+02;
Matches 13; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: Other nucleic acid, Synthetic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Wenderoth, Lind & Ponack
805 Fifteenth Street, N.W., #700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
TITLE OF COMPATIBLE
TITLE OF COMPATIBLE
TITLE OF COMPATIBLE
TITLE OF COMPATIBLE
TITLE OF COMPATIBLE
TITLE OF COMPA
                                                                                                                                                                                                                                                                                                                    Sequence 17, Application US/08241465B Patent No. 5719125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Warren M. Cheek, Jr.
REGISTRATION NUMBER: 33,367
REPRENCE/DOCKET NUMBER:
TELECOMMUNICATION:
TELEPHONE: (202) 371-8850
TELEFAX:
TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: FUJIO SUZUKI
APPLICANT: YUJI HIRAKI
APPLICANT: Kazuhiro TAKAHASHI
APPLICANT: Junko SUZUKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 17: SEQUENCE CHARACTERISTICS: LENGTH: 16 base pairs
                                                                                                                                                                         1239 GCTGGACGTGGCCAT 1253
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-258-152-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      420 CCTAGACAGGAGCAC 434
                                                                                                                                                                                                                   15 GCTGGACATGGCCCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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APPLICANT: Fujio S
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COUNTRY: D.C
COUNTRY: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-241-465B-17
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RESULT 524 US-08-465-485A-16/c 'S Sequence 16, Application US/08465485A 'Parent No. 5831066

CCTAGACTGGATCAC 16

```
APPLICANT: 25EBO, KRISZILMA "..
APPLICANT: QIN, LU
APPLICANT: QIN, LU
TITLE OF INVENTION: METHOD FOR PRODUCTION OF HIGH TITER
TITLE OF INVENTION: OF NAMMALIAN CELLS
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS: 31
CORRESPONDENCE ADDRESS: STREET: 322 LAKESIDE DRIVE
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0
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                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
TITLE OF INVENTION: Regulation of bcl-2 Gene Expression NUMBER OF SEQUENCES: 29
                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 05-UNN 1995
CLASSIFICATION NUMBER: US/08/465,485A
FILING DATE: 05-UNN 1995
CLASSIFICATION: 514
PRIOR APPLICATION NUMBER: US 08/124,256
FILING DATE: 20-SEP-1993
PRIOR APPLICATION NUMBER: US 07/840,716
FILING DATE: 21-FEB-1992
PRIOR APPLICATION NUMBER: US 07/846,692
PRIOR APPLICATION NUMBER: US 07/288,692
PRIOR APPLICATION NUMBER: US 07/288,692
PRIOR APPLICATION NUMBER: US 07/288,692
PRILNG DATE: 22-DEC-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                   ADDRESSE: P.C. STRET: 1755 S. Jefferson Davis Hwy., Suite 400 CITY: Arlington STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: FOTTINGY, Andrew D.

REGISTRATION NUMBER: 34,600
REFERENCE/DOCKET NUMBER: 3335-070-55 CONT
TELEPHONE: (408) 436-2070
TELEPHONE: (408) 436-2075
INPORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TERGITH: 16 base pairs
TERGITH: 16 base pairs
TERGITH: 16 base pairs
TERGITH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18, Application US/08076299A
Parent No. 5834256
GENERAL INFORMATION:
APPLICANT: FINER, MITCHELL H.
                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ROBERTS, MARGO R.
DULL, THOMAS J.
ZSEBO, KRISZTINA M.
QIN, LU
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                                                                                                                                                                                                                                                                                 Floppy disk
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                                                                                                                                                                                                                            ZIP: 22202
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                       U.S.A.
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Gaps
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                                                                                                                                                                                                                                         0.9%; Score 11.8; DB 1; Length 16; 86.7%; Pred. No. 2.3e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
GENERAL INFORMATION:
GOLDENCY, Tritz, Richard
APPLICANT: Goldenberg, Tsvi
APPLICANT: Tritz, Richard
APPLICANT: Tritz, Richard
APPLICANT: Tritz, Richard
APPLICANT: Tritz, Richard
TITLE OF INVENTION: AND/OR PREVENTION OF RESTENOSIS
NUMBER OF SEQUENCES: 35
NUMBER OF SEQUENCES: 35
NUMBER OF SEQUENCES: 35
ADDRESSEE: SEED and BERRY
STREET: 6300 Columbia Center, 701 Fifth Avenue
CITY: Seattle
STATE: Washington
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
TLING DATE: US/08/527,060
FILING TRANDED/DES: US/08/52/6031
FILING CHARACTER ISTICS:
LENGTH: US/08/52/6031
FILING TRANDED/DESS: SINGle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.9%; Score 11.8; DB 1; Best Local Similarity 86.7%; Pred. No. 2.38+02; Matches 13; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 528
US-08-292-620A-1628/c
; Sequence 1628, Application US/08292620A
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 12, Application US/08527060 Patent No. 5834440
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                                                                                                                                                                                                                                                                                                                                                  601 AGCCTGAAGCCTGAC 615
TELEFAX: (206) 002.002.
INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS: LENGTH: 16 base pairs TYPE: nucleic acid TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 GACTCGTCCACGTC 15
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Best Local Similarity
Matches 13; Conserva
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                                                                                                                                                                          linear
                                                                                                                      TYPE: nucleic
STRANDEDNESS:
TOPOLOGY: line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-527-060-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/08527060
Patent No. 583440
GENERAL INFORMATION:
APPLICANT: Goldenberg, Tavi
APPLICANT: Tritz, Richard
TITLE OF INVENTION: RIBOZYME THERAPY FOR THE TREATMENT
TITLE OF INVENTION: AND/OR PREVENTION OF RESTENOSIS
CORRESPONDENCES: 35
CORRESPONDENCES: 35
CORRESPONDENCES: SEED and BERRY
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: THEN PC compatible
COMPUTER: THEN PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PELEASE #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/527,060
FILING DATE: 12-SEP-1995
CLASSIFICATION: 14
ATTORNEY/AGENT INFORMATION:
NAME: MCMAsters, David D.
REGISTRATION NUMBER: 33,963
REGISTRATION NUMBER: 33,963
REFRENCE/POCKET NUMBER: 480124.402C1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELESCOMMUNICATION INFORMATION:
                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/076,299A
FILING DATE: 11-UW-1993
CLASSIFICATION: 435
PRIOR APPLICATION A435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 34,647
RECISTRATION NUMBER: 34,647
RECISTRATION NUMBER: GELL 13.0
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ 10 NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1239 GCTGGACGTGGCCAT 1253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA (genomic) US-08-076-299A-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15 GCTGGACATGGCCCT 1
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Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                       FOSTER CITY
CALIFORNIA
Y: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 6300 ColuCITY: Seattle
STATE: Washington
COUNTRY: USA
ZIP: 98104-7092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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                                                 STATE: C. COUNTRY:
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Sequence 20. Application US/08282197C

| Patent No. 5871730
| GENERAL INFORMATION:
| APPLICANT: Breatinski, Ryszard
| APPLICANT: Dery, Claude V
| APPLICANT: Dery, Claude V
| TITLE OF INVENTION: Thermostable Xylanase DNA, Protein and TITLE OF INVENTION: Methods of Use
| NUMBER OF SECUENCES: 67
| CORRESPONDENCE ADDRESS: 67
| CORRESPONDENCE ADDRESS: 67
| STREET: 1100 New York Ave., NW
| CITY: Washington STATE: DC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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                                                                                                                                                                                                                                                    COMPUTER FRADABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/438,582
FILING DATE: 10-MAY-1995
CLASSIFICATION NUMBER: US 08/258,152
FILING DATE: 10-JUN-1994
PRIOR APPLICATION NUMBER: US 08/076,299
FILING DATE: 11-JUN-93
ATTONEY/AGENT INFORMATION:
NAME: KRUPEN, KAREN IC
REGISTRATION NUMBER: 34,647
REGISTRATION NUMBER: 34,647
REGISTRATION NUMBER: 34,647
REGISTRATION NUMBER: 34,647
REGISTRATION NUMBER: 34,647
REFERENCE/DOCKET NUMBER: 34,647
RELEPERK: 415-358-9600 X131
TELEPERK: 415-358-97392
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                    OF MAMMALIAN CELLS
32
TITLE OF INVENTION: OF MAMMALI
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSE: CELL GENESYS, INC.
STREET: 322 LAKESIDE DRIVE
CITY: FOSTER CITY
STATE: CALIFORNIA
ZONNTRY: USA
ZIP: 94404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1239 GCTGGACGTGGCCAT 1253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 415-349-7392
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15 GCTGGACATGGCCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ð
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 4335
CLASSIFICATION: 4335
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
0.9%; Score 11.8; DB 1;
Best Local Similarity 86.7%; Pred. No. 2.38+02;
Matches 13; Conservative 0; Mismatches 2;
                                                                                                                                APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOXTME TREATMENT OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: NOLECULE-1 (I-CAM-1)
NUMBER OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620A
FILING DATE: August 17, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-438-582-18/c

Sequence 18, Application US/08438582

Patent No. 5858740

GARREAL INFORMATION:

APPLICANT: FINER, MITCHELL H.

APPLICANT: ROBERTS, MARGO R.

APPLICANT: APELO, THOMAS J.

APPLICANT: ZSEBO, KRISZINA M.

APPLICANT: ZSEBO, KRISZINA M.

TITLE OF INVENTION: WETHOD FOR PROFITLE OF INVENTION: VIRUS AND HIGH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               208/149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELERAK: (213) 955-0440
TELEK: 67-310
INFORMATION FOR ESC ID NO. 1628:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
COLOGY: linear
US-08-292-620A-1628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                              Dan T. Stinchcomb
James McSwiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           595 CCCACCAGCCTGAAG 609
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                                                                                                              Sean Sullivan
                                       Susan Grimm
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8, Application US/08817145
; Sequence 8, Application US/08817145
; Sequence 8, Application US/08817145
; Patent No. 6025239
; GENERAL INFORMATION:
    APPLICANT: UTSUMI, Jun
    APPLICANT: TANAKA, Yasuhiko
    APPLICANT: TANAKA, Yasuhiko
    APPLICANT: MATSUI, Mizuo
    APPLICANT: MATSUI, Mizuo
    APPLICANT: TANAKA, Yasuhiko
    APPLICANT: MATSUI, TERRAPEUTIC AGENT FOR OPHTHALMIC
    TITLE OF INVENTION: THERAPEUTIC AGENT FOR OPHTHALMIC
    TITLE OF INVENTION: DISEASES
    ADDRESSEE: Birch, Stewart, Kolasch & Birch, LLP.
    STREET: P.O. Box 747
    CONTRES PALO ON 747
    COUNTY: VA
    COUNTY: VA
    COUNTY: VA
    CONTUTER READABLE FORM:
    MEDIUM TYPE: Floppy disk
    COMPUTER: INP PC COMPATIBLE
    COMPUTER: PLOPS/MS-DOS
    SOFTWARE: PATENTIN PAT:
    COMPUTER: DATEN: O2-UUL-1997
    CLASSIFICATION: 514
    ATTORNEY/AGENT INFORMATION:
    AREFERENCE/DOCKET NUMBER: 760-230P
    TELEPHONE: PON-CONTUTENTION:
    TELEPHONE: PON-CON
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86.7%; Pred. No. 2.3e+02;
tive 0; Mismatches 2;
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DESCRIPTION: /desc = "Synthetic Primer"
                                                                                 TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         625 GACCAGCTCCAGGAG 639
                                                                                                                                                               ANTI-SENSE: NO ORIGINAL SOURCE:
CREATING SOURCE:
CREATING PETUNIA hybrida
US-08-137-024-2
                                                                                                                                                                                                                                                                                                                                                                                                                                387 AGAGGTGGCAGCAAT 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 AGAGGTGACAGAAT 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.99
Best Local Similarity 86.77
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 86.7
Matches 13; Conservative
16 base pairs
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STRANDEDNESS: single
                     TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
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US-08-817-145-8/c
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                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-137-024-2
Sequence 2, Application US/08137024
Fatent No. 6005167
Fatent No. 6005167
FAPLICANT: VAN TUNEN, Adrianus, J.
APPLICANT: VAN DER MEER, Ingrid M.
APPLICANT: WOL, Josephus N.M.
TITLE OF INVENTION: FOR OBTAINING MALE STERILE
TITLE OF INVENTION: PLANTS AND RECOMBINANT DNA FOR TITLE OF INVENTION: USE THEREIN
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 10023
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette 3.50 inch, DS, DD 720
MEDIUM TYPE: KD/720Ko
COMPUTER: IBM PC Compatible 286 SX 12 Mhz
OPERATING SYSTEM: DOS
                                                                                                                                                                     1050.0410000
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/137,024
FILING DATE: 14-OCT-1993
CLASSIFICATION 800
PRIOR APPLICATION BOTA:
APPLICATION NUMBER: PCT/NL92/00075
FILING DATE: 15-APR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 91200910
FILING DATE: 16-APR-1991
ATTORNEY/AGENT INFORMATION:
NAME: MASS, CLIFFCRA, J.
REGISTRATION NUMBER: 30086
APPLICATION NUMBER: US/08/282,197C
FILING DATE: 29-JUL-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: U-9373
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 708-1800
TELEFAX: (212) 246-8959
                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A
REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 1050.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2540
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
ITYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: Ladas & Parry
26 West 61st Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          530 AGGAGCAGCTGGGTG 544
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: New York
STATE: New York
COUNTRY: USA
ZIP: 10023
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 11; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-282-197C-20
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FARSON, DEBORAH A. FRANCISCO OF HIGH TITER MEDIATED TRANSDUCTION INTO AND HIGH EFFICIENCY RETROVIRAL MEDIATED TRANSDUCTION FINITION: OF MAMMALIAN CELLS
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86.7%; Pred. No. 2.3e+02;
tive 0; Mismatches 2;
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: BIBOZYME TREATMENT OF
TITLE OF INVENTION: BIBOZYME TREATMENT OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: NOTECULE-1 (I-CAM-1)
NUMBER OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSER: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: storage
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE:
CLASSIFCATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/ACBAT INFORMATION:
NAME: Warburg, Richard J.
REGESTRATION NUMBER: 208/149
FILING DATE: December 7, 1992
ATTORNEY/ACBAT INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELETSA: 67-3510
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TELETSA: 67-3510
TELETSA: FORMATION FOR SERIES
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COMPUTER READBLE FORM:
MEDIUM TYPE: 35" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18, Application US/09266596; Patent No. 6218187; GENERAL INFORMATION: APPLICANT: FINER, MITCHELL H. APPLICANT: ZEBEO, KRISZTINA M. APPLICANT: COOKE, KEEGAN APPLICANT: FARSON, DEBORAH A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           595 CCCACCAGCCTGAAG 609
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Best Local Similarity 86.7'
...nes 13; Conservative
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EDNESS: single
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ADDRESSEE: CELL GENE
                                                                                                                                                                                                                                             STREET: Suite 4700 CITY: Los Angeles STATE: California COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: M
TITLE OF INVENTION: V
TITLE OF INVENTION: O
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                                                          Sequence 16, Application US/09080285
; Sequence 16, Application US/09080285
; Patent No. 6040181
; GENERAL INFORMATION:
    TITLE OF INVENTION:
    NUMBER OF SEQUENCES: 29
; CORRESPONDENCE ADDRESS:
    ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, ADDRESSEE: P.C.
    STREET: 1755 S. Jefferson Davis Hwy., Suite 400
    CITY: Alington
    STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 11.8; DB 1; Length 16; 86.7%; Pred. No. 2.3e+02; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/080,285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Fortney, Andrew D.
REGISTRATION NUMBER: 34,600
REGISTRATION NUMBER: 3335-070-55 CONT
TELEPHONE: (408) 436-2075
TELEPHONE: (408) 436-2075
INFORMATION FOR SEQ ID NO: 16: SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION UNBER: US 08/465,485
FILING DATE: 05-UN-1995
FILING DATE: 05-UN-1995
FILING APPLICATION NUMBER: US 08/124,256
FILING DATE: 20-SEP-1993
FRICR APPLICATION NUMBER: US 07/840,716
FILING DATE: 21-FEB-1992
FRICR APPLICATION NUMBER: US 07/840,716
FILING DATE: 21-FEB-1992
APPLICATION NUMBER: US 07/288,692
FILING DATE: 22-DEC-1988
FILING DATE: 22-DEC-1988
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Patent No. 6132967
GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA (genomic)
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Best Local Similarity 86.7
Matches 13; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                               STATE: Virginia
COUNTRY: U.S.A.
ZIP: 22202
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                            RESULT 533
US-09-080-285-16/c
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US-09-080-285-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Zwick, Michael G.
APPLICANT: Zwick, Michael G.
APPLICANT: Zwick, Michael G.
APPLICANT: Edington, Brent E.
APPLICANT: McSwiggen, James A.
APPLICANT: McSwiggen, James A.
APPLICANT: Guo, Lining
APPLICANT: Skokut, Thomas A.
APPLICANT: Scott A.
APPLICANT: Folkerts, Octt A.
APPLICANT: Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR TITLE OF INVENTION: IN PLANTS
NUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSED: Lyon & Lyon
ADDRESSED: Lyon & Lyon
ADDRESSED: Lyon & Lyon
                             FILING DATE: 07-Unn-1995
CIASSIFICATION: «Unknown»
PRIOR APPLICATION LAURANDAMS.
APPLICATION AUMBER: 08/238,405
FILING DATE: 05-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Mandel, SaraLynn
REGISTRAATION NUMBER: 31,853
REFERENCE/DOCKET NUMBER: 0c11 5.3
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 358-9600
TELEPHONE: (415) 358-9600
TELEPHONE: (415) 358-0803
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: NUCLEIC acid
STREE CONTONEY: 1 in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a street or in a stree
       APPLICATION NUMBER: US/08/479,737
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear

MOLECULE TYPE: CDNA

SEQUENCE DESCRIPTION: SEQ ID NO: 16:
US-08-479-737-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US/08/679,645
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APPLICATION NUMBER: US/08/679,645
FILING DATE: UJY 12, 1996
CLASSIFICATION: BOD
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 66/001,135
FILING DATE: July 13, 1995
FILING DATE: September 2, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 523, Application US/08679645
Patent No. 6350934
GENERAL INFORMATION:
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US-08-679-645-523
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Roberts, Margo R
Zsebo, Kristina
TITLE OF INVENTION: CHIMERIC CHAINS FOR RECEPTOR ASSOCIATED
SIGNAL TRANSDUCTION PATHWAYS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                ZIP: 94404

COMPUTER READABLE FORM:
MEDIUM TYPEM FOODPOUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/266,596
                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 10955
APPLICATION NUMBER: US 08/258,152
FILING DATE: 10-070-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/076,299
FILING DATE: 11-070-1993
APPLICATION NUMBER: 34,647
REGISTRATION NUMBER: 34,647
REFENCE/DOCKET NUMBER: 34,647
REFENCE/DOCKET NUMBER: CELL 13.3
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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ADDRESSEE: CELL GENESYS, INC.
STREET: 322 Lakeside Drive
CITY: Foster City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CELL 13.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 530
US-08-479-737-16/c
Sequence 16, Application US/08479737
Patent No. 6319494
GENERAL INFORMATION:
APPLICANT: Capon, Daniel J
Weiss, Arthur
Weiss, Arthur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
STREET: 322 LAKESIDE DRIVE
CITY: FOSTER CITY
STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 415-349-7392
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1239 GCTGGACGTGGCCAT 1253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: California
COUNTRY: USA
ZIP: 94404
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
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                                                                                                   COUNTRY:
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Satent No. 03-00-00-00
GENERAL INFORMATION:
APPLICANT: Schlingensiepen, Georg-Ferdinand
APPLICANT: Schlingensiepen, Karl-Hermann
APPLICANT: Schlingensiepen, Karl-Hermann
APPLICANT: Schlingensiepen, Reimar
APPLICANT: Schlingensiepen, Reimar
APPLICANT: Bogdahn, Ulrich
TITLE OF INVENTION: Antisense-oligonucleotides for the treatment of
TITLE OF INVENTION: Immuno-suppressive effect of transforming-growth-factor beta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                    Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Reed, John
TITLE OF INVENTION: Regulation of BCL-2 Gene Expression
FILE REFERENCE: 10412-024
CURRENT APPLICATION NUMBER: US/09/724,426
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PatentIn version 3.0
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                  0.9%; Score 11.8; DB 1;
86.7%; Pred. No. 2.3e+02;
tive 0; Mismatches 2;
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Jacobson, Price, Holman & Stern
STREET: 400 Seventh St. N.W.
CITY: Washington D.C
COUNTRY: U.S.A.
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APPLICATION NUMBER: US/08/535,249
FILING DATE:
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-724-426-16/c
; Sequence 16, Application US/09724426
; Patent No. 6414134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 97, Application US/08535249 Patent No. 6455689
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                 13; Conservative
                                                                                           single
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                         Best Local Similarity
                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                    STRANDEDNESS:
                                                                                                           TOPOLOGY: 11: MOLECULE TYPE:
                                                                                                                                                          US-08-475-442A-16
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LENGTH: 16
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Sequence 16, Application US/08475442A

Patent No. 6407221

GENERAL INFORMATION:

APPLICANT: CAPON, DANIEL J

APPLICANT: REVING, BRIAN A

APPLICANT: REVING, BRIAN A

APPLICANT: RESEO, KRISZTINA

APPLICANT: RESEO, KRISZTINA

TITLE OF INVENTION: RECEPTOR-ASSOCIATED SIGNAL TRANSDUCTION PATHWAYS

TITLE OF INVENTION: RECEPTOR-ASSOCIATED SIGNAL TRANSDUCTION PATHWAYS

NUMBER OF SEQUENCES: 51

CORRESPONDENCE ADDRESS:

ADDRESSEE: CELL GENESYS, INC.
                                                                                                                                                                                                                                                                                                                                                                                                           .
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/238,405
FILING DATE: 05-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/988,194
FILING DATE: 09-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/627,643
FILING DATE: 14-DEC-1990
PRIOR APPLICATION DATA:
                  NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TELEPHONE: (213) 489-1600
TELEPA: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 523:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          UMBER: US/08/475,442A
06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/US91/09431
FILING DATE: 12-DEC-1991
ATTORNEY/AGENT INFORMATION:
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TELEPHONE: (415)358-9600x131
TELEFAX: (415)349-7392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              34,647
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CITY: FOSTER CITY
STATE: CALIFORNIA
COUNTRY: USA
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  ATTORNEY/AGENT INFORMATION:
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REGISTRATION NUMBER:
                                                                                                                                                                                                                         LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 06-JU
                                                                                                                                                                                                                                                                                             linear
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Matches 10; Conserv
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COUNTRY: USI
ZIP: 94404
                                                                                                                                                                                                                                                                                                                US-08-679-645-523
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TELEFAX: 415-349-7392
INFORMATION FOR SEQ ID NO: 18:
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LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1239 GCTGGACGTGGCCAT 1253
                                                                                                                                                                                                                                                                                                                                                          ZIP: 94404
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                 STATE: CALIFORNIA COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                              APPLICANT: FINER,
                                                                                       ZSEBO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 13; Conserv
                          GENERAL INFORMATION:
      Patent No. 6506604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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APPLICANT: Sparks, Andrew
APPLICANT: Sparks, Andrew
APPLICANT: Sparks, Andrew
APPLICANT: Sparks, Andrew
APPLICANT: Minzler, Kenneth
APPLICANT: Wogelstein, Bert
TITLE OF INVENTION: Serial analysis of transcript expression
TITLE OF INVENTION: Using long tags
FILE REFERENCE: 001107: 00172
CURRENT APPLICATION NUMBER: US/09/916,228
CURRENT APPLICATION NUMBER: 60/227
FRIOR FILING DATE: 2000-07-28
FRIOR FILING DATE: 2000-09-18
FRIOR FILING DATE: 2000-09-18
NUMBER OF SEQ ID NOS: 30
SOFTWARE: FastSEQ for Windows Version 4.0
IENGTH: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.9%; Score 11.8; DB 1; Length 16; 86.7%; Pred. No. 2.3e+02; tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
0.9%; Score 11.8; DB 1;
Best Local Similarity 86.7%; Pred. No. 2.3e+02;
Matches 13; Conservative 0; Mismatches 2;
PRIOR APPLICATION DATA:

APPLICATION NUMBER: EP 93 107 089.0
FILING DATE: 30-APR-1993
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: EP 93 107 849.7
FILING DATE: 13-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Player, William E.
REGISTRATION NUMBER: 1409
REFERENCE/DOCKET NUMBER: 10577/P58418
TELEPRANT (202) 638-666
TELEPRAN: RCA 24859 1DRA UR
INFORMATION FOR SEQ 1D NO: 97:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: mucleic acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: tag or tag concatamer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 14, Application US/09916228
Patent No. 6498013
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 18, Application US/09944411
                                                                                                                                                                                                                                                                                                                                                                                             unknown
PE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1023 GTGCCAAAGTGCAGC 1037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               190 GCCGCCCACCCGGAC 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GTACAAAGTGCAGC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GCCGCTCTCCCGGAC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: unkr.
MOLECULE TYPE: L
ANTI-SENSE: YES
US-08-535-249-97
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US-09-944-411-18/c
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US-09-916-228-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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APPLICANT: ENEW ANTOREL H.

SERIO, MESSATINA M.

DILL, THOMAS J.

SERIO, MESSATINA M.

SERIO, SERIOR MESSATINA M.

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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (BPO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Nixon and Vanderhye
STREET: 1100 No. 6555654th Glebe Road, Eighth Floor
STREET: Azlington
STATE: Virginia
COUNTRY: US
CONTRY: US
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                       0.9%; Score 11.8; DB 1; Length 16; 86.7%; Pred. No. 2.38+02; Live 0; Mismatches 2; Indels
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Kawaguchi, Yoshihiko
Merziman, Tony R
Merzker, Michael L
TITLE OF INVENTION: No. 6555654el LDL-Receptor
NUMBER OF SEQUENCES: 455
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/402,923A
FILING DATE: 14-Feb-2001
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB98/01102
FILING DATE: 15-APR-1998
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
APPLICATION NUMBER: US 67/48,740
FILING DATE: 05-JUN-1997
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REFERENCE/DOCKET NUMBER: 620-81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 420, Application US/09402923A
Patent No. 6555654
GENERAL INFORMATION:
Hess, John W
Caskey, Charles T
Cox, Roger D
Gerhold, David
Hammond, Holly
APPLICATION NUMBER: US 60/048,740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)816-4091
TELEFAX: (703)816-4100
          ### FILING DATE: 05-JUN-1997
# ATTORNEY/AGENT INFORMATION:
# ANAME: B.U.Sadoff
# REGISTRATION NUMBER: 36,663
# REFERENCE/DOCKET NUMBER: 620-3
# TELECOMMUNICATION INFORMATION:
# TELEPHONE: (703)816-4100
# INFORMATION FOR EQO ID NO: 420:
# SEQUENCE CHARACTERISTICS:
# LENGTH: 16 base pairs
# TYPE: nucleic acid
# TYPE: nucleic acid
# TYPE: nucleic acid
# TOPOLOGY: linear
# US-09-060-299-420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION: NAME: B.J.Sadoff
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 420
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Best Local Similarity 86.7
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Nixon and Vanderhye
STREET: 1100 No. 6545137th Glebe Road, Eighth Floor
CITY: Arlington
STATE: Visginia
COUNTRY: US
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: EIB PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,299
FILING DATE: 15-APR-1998
CLASSIFICANTON: 435
                                                          APPLICANT: Kawaguchi, Yoshihiko
APPLICANT: Merriman, Tony R
APPLICANT: Metzker, Michael L
TITLE OF INVENTION: No. 6545137el Receptor
NUMBER OF SEQUENCES: 455
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 420, Application US/09060299; Patent No. 6545137; GENERAL INFORMATION: APPLICANT: Hess, John WAPPLICANT: Caskey, Charles TAPPLICANT: Cox, Roger DAPPLICANT: Gerhold, David APPLICANT: Hammond, Holly APPLICANT: Hammond, Holly APPLICANT: Hew, Patricia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                720 CCAGCAGCAGGGGCC 735
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STRANDEDNESS: single
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US-08-754-477A-38
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                      PCT-US96-00331-19/c
; Sequence 19, Application PC/TUS9600331
GENERAL INFORMATION:
APPLICANT: GENTA INCORPORATED
TITLE OF INVENTION: METHODS AND COMPOSITION FOR
TITLE OF INVENTION: TREATING TUMOR CELLS
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/00331
FILING DATE: 10 JANUARY 1996
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/371,001
FILING DATE: 10 JANUARY 1995
ATTORNEY AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 218/068-PCT
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Other Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
       SOFTWARE: PatentIn version 3.0 SEQ ID NO 5661
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: (7.3510
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                           300 TGCTGTGGGGGCTGC 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        487 CGAGACGGTGTGCAG 501
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                                                                       TYPE: RNA
ORGANISM: Homo sapiens
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STATE: California
COUNTRY: U.S.A.
                                                                                                                                                              Query Match
Best Local Similarity
Matches 9; Conserva
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Best Local Similarity
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ANTI-SENSE: Yes
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                                                                                                                    US-09-371-772B-5661
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                                                    LENGTH
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APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: MABBR: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Structon, Jim
APPLICANT: Structomb, Dan
APPLICANT: Structcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: MAHBER: US/09/371,772B
CURRENT FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 06/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR RILING DATE: 1996-01-08
SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEQ ID NO 5660
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                                                                                                                                                                                      Length 16;
                                                                                                                                                                                                                                 2; Indels
                                                                                                                                                                                 Score 11.8; DB 1;
Pred. No. 2.3e+02;
0; Mismatches 2;
                     LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 420:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 5661, Application US/09371772B Patent No. 6566127 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 5660, Application US/09371772B
Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                 0.9%;
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Best Local Similarity 86.7%
Matches 13; Conservative
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Best Local Similarity
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US-09-371-772B-5660
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US-09-402-923A-420
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1300 I Street N.W., Suite 700
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Job time : 17 secs
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                      Washington
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                                         GENERAL INFORMATION:
APPLICANT: Levitt, Roy C.
APPLICANT: Maloy, W. Lee
APPLICANT: Kari, U. Prasad
APPLICANT: Nicolaides, Nicholas C.
TITLE OF INVENTION: Asthma Associated Factors As Targets For
TITLE OF INVENTION: Treating Atopic Allergies Including Asthma And Related
NUMBER OF SEQUENCES: 41
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Levitt, Roy C.
APPLICANT: Levitt, Roy C.
APPLICANT: Maloy, W. Lee
APPLICANT: Maloy, W. Lee
APPLICANT: Micolaides, Nicholas C.
TITLE OF INVENTION: Asthma Associated Factors As Targets For
TITLE OF INVENTION: Disorders
TITLE OF INVENTION: Disorders
NUMBER OF SEQUENCES: 41
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
ADDRESSEE: Dunner L.L.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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                                                                                                                                                                                                                                                                                        ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & ADDRESSEE: Dunner L.L.P.
STREEF: 1300 I Street N.W., Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATE: PATEN: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/702,105A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.9%; Score 11.6; DB 1; Best Local Similarity 77.8%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Fordis, Jean B.
REGISTRATION NUMBER: 32984
REFERENCE/DOCKET NUMBER: 05387.0056-01000
TELECOMMUNICATION INFORMATION:
TELEFHONE: (202) 408-4400
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
PAPLICATION NUMBER: US/08/874,503
FILING DATE: 13-UIN-1997
ATTORNEY/AGENT INFORMATION:
       Sequence 33, Application US/08702105A
Patent No. 5908839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-702-110A-33; Sequence 33, Application US/08702110A; Patent No. 6037149; GENERAL INPORMATION;
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MOLECULE TYPE: DNA (genomic)
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                STATE: D.C. COUNTRY: USA
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                                                                                                                                               COMPUTER: LIBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
APPLICATION NUMBER: US/08/702,110A
FILING DATE:
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Pred. No. 3.2e+02;
0; Mismatches 4;
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ER: 05387.0056-01000
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CLASSIFICATION:

PRIOR APPLICATION DATA:

APPLICATION UNDER:

TILING DATE: 13-UN-1997

ATTORNEY/AGENT INFORMATION:

NAME: Fordis, Jean B.

REGISTRATION NUMBER: 05387.0056-

REGISTRATION NUMBER: 05387.0056-

TELEFAN: (202) 408-4000

TELEFAN: (202) 408-4000

TELEFAN: (202) 408-4000

TELEFAN: (202) 408-400

TELEFAN: (202) 408-4000

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STATE: D.C.
COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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Best Local Similarity 77.8%;
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